

REMARKS

Favorable reconsideration is respectfully requested in view of the foregoing amendments and the following remarks.

Initially, Applicants and their undersigned representative wish to express their sincere appreciation to the Examiners for their courtesy and helpful comments made during the personal interview.

Claims 5-7 have been amended to more particularly define the present invention. Further, claim 20 has been amended to correct an inadvertent error. Support for the claim amendments is readily apparent from the teachings of the specification and the original claims.

With regard to the rejection of claims 1, 3, 5-7 and 10-30 under 35 USC § 102(a) as being anticipated by Kumagai et al. as set forth in item 8 of the Official Action, this rejection has been overcome by the filing of the verified English translation of the certified priority document enclosed herewith.

With regard to the rejection of claims 1, 3, 18 and 19 under 35 USC § 112, second paragraph, this rejection is deemed to be untenable and is thus respectfully traversed. The Examiners conceded during the personal interview that the term “*comprising*” is a well recognized term under U.S. practice and is thus, not vague or indefinite under 35 USC § 112, second paragraph. Although the term renders the claims to open interpretation, the breadth of a claim is not to be equated with indefiniteness. *In Re Miller*, 441 Fed. 2nd 689, 169 USPQ597 (CCPA 1971). If the scope of the subject matter set forth by the claims is clear, and if Applicants

have not otherwise indicated that they intended the invention to be of a scope different from that defined in the claims, then the claims comply with 35 USC § 112, second paragraph.

Since, in this case, the scope of the claims is clear and since it precisely correspond to that which is intended by the Applicants (*to encompass all cDNAs and polynucleotides which comprise the amino acid sequence of SEQ ID Nos. 1 or 2*), this rejection can not be sustained and should be withdrawn.

Attached hereto is a marked-up version of the changes made to the claims by the current amendment. The attached page is captioned "VERSION WITH MARKINGS TO SHOW CHANGES MADE."


In view of the foregoing amendments and remarks, it is respectfully submitted that the Application is now in condition for allowance. Such action is thus respectfully solicited.

If, however, the Examiner has any suggestions for expediting allowance of the application or believes that direct communication with Applicants' attorney will advance the prosecution of this case, the Examiner is invited to contact the undersigned at the telephone number below.

Respectfully submitted,

Kenichi ARAI et al.

By: _____


Lee Cheng
Registration No. 40,949
Attorney for Applicants

LC/gtg
Washington, D.C. 20006-1021
Telephone (202) 721-8200
Facsimile (202) 721-8250
February 28, 2003

VERSION WITH MARKINGS TO SHOW CHANGES MADE

The claims have been amended as follows.

5. **(Twice Amended)** An isolated [human gene] cDNA encoding the human protein of claim 1 or 3.

6. **(Twice Amended)** [A] The cDNA [of the human gene] of claim 5, which comprises a nucleotide sequence of SEQ ID NO: 3.

7. **(Twice Amended)** [A] The cDNA [of the human gene] of claim 5, which comprises a nucleotide sequence of SEQ ID NO: 4.

20. **(Amended)** An isolated polynucleotide encoding the polypeptide of claim [1] 18 or [3] 19.

VERIFICATION OF TRANSLATION

I, Mariko Ogura

of 6th Fl., Kantetsu Tsukuba-Science-City Bldg. 1-1-1, Oroshi-machi, Tsuchiura, Ibaraki,
JAPAN

declare as follows:

1. That I am well acquainted with both the English and Japanese languages, and
2. That the attached document is a true and correct translation made by me to the best of my knowledge and belief of:-
 - (a) Japanese Patent Application No. Hei 10-311408
Entitled: "HUMAN H37 PROTEIN AND cDNA ENCODING THE PROTEIN"
Filed on October 30, 1998

January 31, 2003
(Date)

Mariko Ogura
Mariko Ogura
(Signature of Translator)

【Name of Document】 APPLICATION FOR PATENT

【Identification Number】 NP98282-YS

【Filing Date】 October 30, 1998

【Address】 Esq. Commissioner of the Patent Office

【International Patent Classification】 C07K 13/00

**【Title of the Invention】 HUMAN H37 PROTEIN AND cDNA ENCODING
THE PROTEIN**

【Number of Claims】 17

【Inventor】

【Address】 9-6-206, Meguro 1-chome, Meguro-ku, Tokyo

【Name】 Kenichi Arai

【Inventor】

【Address】 Shanboursu-Mita 620, Mita 5-7-8, Minato-ku, Tokyo

【Name】 Hisao Masai

【Applicant】

【Identification Number】 396020800

【Name or Appellation】 JAPAN SCIENCE AND TECHNOLOGY CORPORATION

【Applicant】

【Address】 9-6-206, Meguro 1-chome, Meguro-ku, Tokyo

【Name or Appellation】 Kenichi Arai

【Applicant】

【Address】 Shanboursu-Mita 620, Mita 5-7-8, Minato-ku, Tokyo

【Name or Appellation】 Hisao Masai

【Attorney】

【Identification Number】 100093230

【Patent Attorney】

【Name or Appellation】 Toshio Nishizawa

【Telephone Number】 03-5454-7191

【Payment】

【Registration Number】 009911

【Amount】 21000 yen

【List of Attached Documents】

【Name of Document】 Specification 1

【Name of Document】 Drawings 1

【Name of Document】 Abstract 1

【Proof】 Necessary

[Document Name] Specification

[Title of the Invention] Human H37 protein and cDNA encoding the Protein

[Claims]

5 [Claim 1] Human H37 protein having an amino acid sequence of SEQ ID NO: 1.

[Claim 2] Human H37 protein having an amino acid sequence where one or more amino acid residue(s) in the amino acid sequence of SEQ ID NO: 1 is/are deleted therefrom, substituted therefor or added
10 thereto.

[Claim 3] Human H37 protein having an amino add sequence of SEQ ID NO: 2.

[Claim 4] Human H37 protein having an amino acid sequence where one or more amino acid residue(s) in the amino acid sequence of SEQ ID NO: 2 is/are deleted therefrom, substituted therefor or added
15 thereto.

[Claim 5] A human gene encoding the human H37 protein of claim 1 or 2.

[Claim 6] A cDNA of the human gene of claim 5, which has a base sequence of SEQ ID NO: 3 or a base sequence where one or more base(s) in the base sequence of SEQ ID NO: 3 is/ are deleted therefrom, substituted therefor or added thereto.
20

[Claim 7] A cDNA of the human gene of claim 5, which has a base sequence of SEQ ID NO: 4 or a base sequence where one or more base(s) in the base sequence of SEQ ID NO: 4 is/are deleted therefrom, substituted therefor or added thereto.
25

[Claim 8] A DNA fragment comprising a partial sequence of cDNA of claim 6.

[Claim 9] A DNA fragment comprising a partial sequence of cDNA of claim 7.
30

[Claim 10] A recombinant vector having the cDNA of claim 6.

[Claim 11] A recombinant vector having the cDNA of claim 7.

[Claim 12] An antibody against the human H37 protein of claim 1 or 2.

35 [Claim 13] An antibody against the human H37 protein of claim 3 or 4.

[Claim 14] A method for promoting the proliferation of cells, which comprises introducing the cDNA of claim 6 or the DNA fragment of claim 8 together with expression-regulatory sequences into the cells.

5 [Claim 15] A method for promoting the proliferation of cells, which comprises introducing the cDNA of claim 7 or the DNA fragment of claim 9 together with an expression regulatory sequence into the cells.

[Claim 16] A method for suppressing the proliferation of cells,
10 which comprises introducing the antibody of claim 10 into the cells.

[Claim 17] A method for suppressing the proliferation of cells, which comprises introducing the antibody of claim 11 into the cells.

[Detailed Description of the Invention]

[0001]

15 [Technical Field of Industrial Application]

This application relates to human H37 protein and to cDNA encoding the protein. More particularly, this application relates to human H37 protein that is an activity-controlling subunit for the protein Cdc7 controlling the replication of human cells; to human
20 gene encoding the protein; to an antibody to the H37 protein; and to a method for controlling the proliferation of human cells using such genetic engineering material and antibody.

[0002]

[Prior Art]

25 Proliferation of cells is initiated when a liquid factor called a growth factor is bound to the receptor on the cell surface and a signal for proliferation is transmitted into the cell. Accordingly, for an artificial induction of proliferation of incubated cells, methods where an excessive amount of a growth factor is added to a
30 cell medium, where a receptor which is not inherently owned by the said cell is expressed on the cell surface and a factor which is specific for the receptor is added to a medium, etc. have been carried out. Further, in suppressing the cell proliferation, methods where competing molecule, antagonist or the like to the receptor protein
35 is added to the medium to suppress the binding of the receptor to the growth factor, etc. have been carried out.

[0003]

On the other hand, in the case of the cell where a proliferation signal is issued by binding of the receptor to the growth factor, a cycle in which its genomic DNA is replicated, uniformly distributed to daughter cells and then divided is repeated. Such a cycle is called "cell cycle" especially for eukaryotes. The cell cycle is basically classified into four phases. Thus, S phase when chromosomal DNA is replicated; M phase when the replicated chromosome is divided by a spindle body and then cytoplasm is divided; G1 phase which is a period from M phase finishes until S phase begins; and G2 phase which is a period from S phase finishes until M phase starts. Transition from G1 phase to S phase is particularly strictly controlled and DNA replication takes place only once in S phase.

[0004]

It has been confirmed from the studies in yeast and higher eukaryotic cells that cyclin-dependent kinase plays a critical role in the cell cycle progression (Nature 292:558-560, 1981; Cell 66:731-742, 1991; Nature 349:338-393, 1991; Science 257:1958-1961, 1992; Bioassays 17:471, 1995). Further, from a genetic analysis in yeast, it has been clarified that another serine/threonine kinase plays an essential role in the initial stage of S phase (G1/S transition). Thus, characterization of cdc7 mutation which was isolated as one of the cell division cycle mutants (J. Mol. Biol. 59:183-194, 1971), it revealed that the Cdc7 protein kinase functions immediately prior to chromosomal replication and that, during S phase, that is necessary for activation of origins (Mol. Cell Biol. 6:1590-1598, 1986; Genes Dev. 15:480-490, 1998; Genes Dev. 15:491-501, 1998). It has been also clarified that the Cdc7 kinase activity is dependent upon the presence of a regulatory subunit, Dbf4 (Genetics 131:21-29, 1992; Mol. Cell. Biol. 13:2899-2908, 1993). Expression of Dbf4 is periodic and is regulated by both at the transcriptional and post-translational levels (Exp. Cell Res. 180:419-428, 1989). The increase in Cdc7 kinase activity at the G1/S boundary is at least accounted for by the elevated expression of Dbf4 in late G1 phase (Mol. Cell. Biol. 13:2899-2908, 1994; Exp. Cell Res. 180:419-428, 1989). In addition, since Dbf4 interacts with replication origins

in vivo (Science 265:1243-1246, 1994), it has been suggested that the Cdc7 may trigger S phase by directly activating the replication initiation complex assembled at the origins.

[0005]

5 In addition, the inventors of this application had already isolated kinases related to yeast Cdc7 from *Schizosaccharomyces pombe*, *Xenopus*, mouse and human (J. Biol. Chem. 273:23248-23257, 1998; EMBO J. 16:4340-4351, 1997; EMBO J. 14:3094-3104, 1995), and pointed out that eukaryotic chromosomal replication is regulated by
10 a conserved mechanism involving this family of kinase.

[0006]

[Problems to Be Solved by the Invention]

From the findings in yeast and higher eukaryotes as mentioned above, it is expected that an artificial control of the cell
15 proliferation is possible by means of the regulation of the Cdc7 kinase activity in cells, which is entirely other means than conventional methods using operation of growth factor/receptor binding.

[0007]

However, the inventors had also found that the putative human
20 homolog of Cdc7, huCdc7, possesses only a very low level of kinase activity when overexpressed in mammalian cells while a baculovirus expressed form of huCdc7 is inactive.

Under such circumstances, the inventors of this application have investigated a human cDNA library with a presumption of the
25 presence of a regulatory subunit for human Cdc7, succeeded in isolating the cDNA encoding a novel protein which regulates the kinase activity by binding to huCdc7 and named the said protein encoded in this cDNA as H37 protein.

[0008]

30 An object of the invention is to provide the novel protein obtained by the inventors in an industrially applicable form.

Another object of the invention is to provide a human gene encoding the protein and to provide a material for genetic operation including cDNA derived from the gene, an antibody to the protein.

35 [0009]

Still another object of the Invention is to provide a method

for the artificial control of proliferation of human cells using the above material for genetic operation.

[0010]

[Means to Solve the Problems]

5 This application provides human H37 protein having an amino acid sequence of SEQ ID NO: 1 or NO: 2.

 This application also provides human H37 protein having an amino acid sequence where one or more amino acid residue(s) in the amino acid sequence of SEQ ID NO: 1 or NO: 2 is/are deleted therefrom,
10 substituted therefor or added thereto.

[0011]

 This application further provides a human gene encoding the above-mentioned human H37 protein; cDNA of said human gene which has a base sequence of SEQ ID NO: 3 or NO: 4; and DNA fragment comprising
15 a partial sequence of those cDNAs.

 This application furthermore provides a recombinant vector having the above-mentioned cDNA and an antibody against the human H37 protein.

[0012]

20 This application still further provides a method for promoting the proliferation of cells, which comprises introducing the above-mentioned cDNA or the DNA fragment together with expression regulatory sequence into cell, and a method for suppressing the proliferation of cells, which comprises introducing the
25 above-mentioned antibody into the cell.

 The present invention will be explained in detail below with respect to Mode for Carrying out the Invention.

[0013]

[Mode for Carrying out the Invention]

30 The human H37 protein of this invention has the amino acid sequence of SEQ ID NO: 1 and is a protein molecule encoded in the sequence region from 518th to 2541st of cDNA whose base sequence is shown in SEQ ID NO: 3. The H37 protein of this invention also has the amino acid sequence of SEQ ID NO: 2 and is a protein encoded in
35 the sequence region from 518th to 1222nd of cDNA whose base sequence is shown in SEQ ID NO: 4. SEQ ID NO: 3 and NO: 4 are cDNAs derived

from mRNA transcribed from the same genomic gene but the cDNA of SEQ ID NO: 4 is in a different splicing form from SEQ ID NO: 3, and there is a deletion from 1199th to 1259th of SEQ ID NO: 3.

[0014]

5 Those H37 proteins can be prepared by known methods such as isolation from human organs and cell lines, chemical synthesis of peptide based upon the amino acid sequence provided by this invention, and a recombinant DNA technique using cDNA fragments provided by this invention. For example, in case the H37 protein is prepared by means
10 of a recombinant DNA technique, RNA is prepared from a vector having the cDNA fragment of this invention by an in vitro transcription and then an in vitro translation is carried out using the above as a template whereby an expression in vitro is possible. In addition, when the translational region is recombined to a suitable vector by
15 a known method, it is possible to express the H37 protein encoded in cDNA in a large quantity in Escherichia coli, Bacillus subtilis, yeast, insect cells, animal cells, etc.

[0015]

When the human H37 protein of this invention is expressed in
20 a microorganism such as E. coli, the translational region of cDNA of this invention is inserted into an expression vector having, for example, promoter, ribosome binding site, cDNA cloning site, terminator, origin replicable in microorganism, etc. whereupon a recombinant expression vector is prepared, then host cell is
25 transformed using the said expression vector and the resulting transformant is incubated whereupon H37 protein encoded in cDNA can be produced in a microorganism in a large quantity. Alternatively, it can be expressed as a fusion protein with other protein. The resulting fusion protein is cleaved by an appropriate protease
30 whereupon only the protein portion encoded in cDNA can be obtained.

[0016]

When the human H37 protein of this invention is expressed in animal cells, the translational region of cDNA of this invention is recombined with expression vector for animal cells having, for example,
35 promoter, poly(A) addition site, and splicing site, and then introduced into the animal cells whereupon the H37 protein of this

invention can be expressed in animal cells.

The human H37 protein obtained by the method as mentioned above can be used as an antigen for the preparation of antibody for suppressing the proliferation of cells through inhibition of the
5 kinase activity of huCdc7.

[0017]

In addition, as will be confirmed by the Examples later, the human H37 protein of this invention has almost no similarity in terms of structure to cyclins which have been elucidated up to now, but
10 it may be regarded as a cyclin-like associating factor for huCdc7 kinase in such respects that its expression is regulated by a cell cycle and that, as a result of binding to an huCdc7 catalytic subunit, its kinase activity can be activated. Accordingly, since the H37 protein is believed to be a very important target factor in a signal
15 transduction pathway for the cell proliferation induced by a growth factor, elucidation of how the expression of the H37 protein or activity thereof is regulated by the signal of cell cycle in G1/S phases is expected to provide a big and novel finding for clarifying the molecular mechanism of cell cycle control of cell replication
20 in animal cells.

[0018]

The human H37 protein of the invention contains peptide fragments (five or more amino acid residues) containing any partial amino acid sequence of SEQ ID NO: 1 or NO: 2. Such peptide fragments
25 can be also used as an antigen for the preparation of antibody.

The gene of the invention is a human gene encoding the above-mentioned human H37 protein and can be isolated from known genomic libraries using, for example, the cDNA of this invention or a partial sequence thereof as a probe.

30 [0019]

The cDNA of the invention is characterized in having a base sequence of SEQ ID NO: 3 or NO: 4, and can be cloned, for example, from cDNA libraries derived from human cells. cDNA is synthesized using poly(A)⁺ RNA extracted from human cells as a template. The human
35 cells may be either those excised from human body by a surgical operation or cell lines. The cDNA can be prepared by known methods

such as Okayama-Berg method (Okayama, H and Berg, P., Mol. Cell. Biol., 2:161-170, 1982), Gubler-Hoffmann method (Gubler, U. and Hoffman, J. Gene, 25:263-269, 1983) and Capping method (Kato, S. et al., Gene, 150:243-250, 1994).

5 [0020]

The human H37 protein of the invention is expressed in any tissue except brain and kidney and, therefore, when the human cDNA library prepared from human cells is screened using an oligonucleotide probe synthesized based on the base sequence of cDNA in SEQ ID NO: 3 or
10 NO: 4, it is possible to easily prepare the same clone of the invention. It is also possible that the desired cDNA is synthesized by a polymerase chain reaction (PCR) using such an oligonucleotide as a primer.

[0021]

15 In general, polymorphism due to the difference among individuals is frequently noted in human gene. Accordingly, cDNA of SEQ ID NO: 3 or NO: 4 where one or more nucleotide(s) is/are added thereto and/or deleted therefrom and/or other nucleotide(s) is/are substituted therefor is also covered by the invention.

20 Similarly, the protein where one or more nucleotide(s) is/are added thereto and/or deleted therefrom and/or other nucleotide(s) is/are substituted therefor as a result of those modifications is also covered by the invention so far as it has an activity of the protein having an amino acid sequence of SEQ ID NO: 1 or NO: 2. Further,
25 variant protein where one or more nucleotide(s) is/are added thereto and/or deleted therefrom and/or other nucleotide(s) is/are substituted by an artificial means is covered by this invention as well.

[0022]

30 The DNA fragment of the invention covers a cDNA fragment (10 bp or more) containing any partial base sequence of SEQ ID NO: 3 or NO: 4, or a cDNA fragment comprising an antisense strand thereof.

Antibody against human H37 protein of the invention can be prepared as a polyclonal antibody or a monoclonal antibody by a known
35 method using protein per se or a partial peptide thereof.

[0023]

A method for promoting the cell proliferation according to the invention is carried out by such a manner that a recombinant DNA comprising the cDNA having a base sequence of SEQ ID NO: 3 or NO: 4, or comprising a partial sequence thereof (such as a DNA fragment encoding 419 amino acid sequence region at the C terminal as shown in Example 3) and an expression-regulatory sequence (promoter and/or enhancer sequence for animal cells) is introduced into animal cells whereby the H37 protein having an amino sequence of SEQ ID NO: 1 or NO: 2 is excessively expressed in cell nuclei. Introduction of the recombinant DNA into cells can be carried out by a known method such as a calcium phosphate method, a method where ribosome and erythrocyte ghost are used, an electroporation method, a method where retrovirus or adenovirus is used as a vector and a microinjection method using a glass pipette. Promotion of cell proliferation as such is useful, for example, in a large-scale preparation of the stem cell that is useful for the therapy of human diseases. Stem cells can be differentiated into other kinds of cells such as blood stem cell and nerve stem cell, and are able to produce a large number of cells of different cell type that constitute the human body. Therefore, transplantation of stem cell in the diseases such as leukemia is a very important therapeutic means. However, since no liquid factor for self-proliferation of human stem cells without differentiation has been identified yet, the preparation of stem cells in an amount sufficient for the therapy has not been easy. According to the method of this invention, it could be possible that, as a result of manipulating the proliferation program in stem cells, the stem cells can be unlimitedly self-replicated and self-proliferated in vitro. In addition, such a promotion of cell proliferation in vitro could be also useful for the preparation of a large quantity of cells for introduction of gene for genetic therapy by ex vivo means.

[0024]

The method for the suppression of cell proliferation according to the invention is carried out by injecting the above-mentioned antibody into cells. Alternatively, it may be carried out by inhibiting the expression of the cell-intrinsic H37 protein gene as well. An example thereof is a method where DNA encoding the ribozyme

sequence or the antisense sequence to the transcript of the gene is introduced into the cells. Suppression of the cell proliferation as such is expected to provide a novel means for suppressing an excessive proliferation of cancer cells for example.

5 [0025]

[Examples]

The invention will now be further illustrated by way of the following examples although the invention is not limited to those examples.

10 Example 1

cDNA library of HeLa cell was prepared using a pGAD-GH vector and each vector was introduced into yeast strain CG1945 that has a recombinant plasmid with huCdc7 fused to DNA-binding domain of Gal4. As a result of a screening of about 3×10^5 cells of transformed yeast, 15 five clones where β -galactosidase was positive were obtained. DNA base sequence of the insert was determined and the database was searched whereupon all of them were novel cDNA. Among them, three clones were same having a base sequence of SEQ ID NO: 1. This cDNA was named H37. Other two were single clones and were named H1 and 20 H18, respectively.

[0026]

The interaction of huCdc7 with protein encoded in those positive clones was further investigated using a production increase system in animal cells. Thus, each expression plasmid of H1, H18 and H37 25 tagged with myc epitope was transfected to animal cell Cos7 together with a full-length huCdc7 expression plasmid. The result is as shown in Fig. 1. Although the H37 protein was coimmunoprecipitated with an antibody against huCdc7, H1 and H18 proteins were not coprecipitated (cf. lanes 2-4 of the upper panel, Fig. 1). On the 30 contrary, huCdc7 was coimmunoprecipitated by using anti-myc antibody alone in the case of the cells where myc-tagged H37 is co-expressed (cf. lane 4 of the lower lane, Fig. 1). From those results, it has been confirmed that only H37 cDNA encodes a protein efficiently interacting with huCdc7.

35 [0027]

Then, in order to investigate the intrinsic H37 protein, each

antibody against N- or C-terminal regions of H37 (anti-H37N antibody or anti-H37C antibody) was prepared. Further, antibody against C-terminal oligopeptide of H37 (anti-H37Cpep antibody) and antibody against C-terminal oligopeptide of huCdc7 (anti-huCdc7Cpep antibody) were also prepared. After that, association of each antibody with intrinsic huCdc7 and H37 in cells was measured. The result is as shown in Fig. 2. Thus, all of the antibodies to H37 specifically react with the myc-tagged H37 protein of 90 kDa expressed in Cos7 cells (cf. lanes 1-4, Fig. 2). Complex prepared from human CEM cells was able to be coprecipitated using an affinity-purified anti-peptide antibody. It was confirmed by an immunoblotting using anti-huCdc7Cpep antibody that all of huCdc7 and H37 immunoprecipitates contained huCdc7 (lanes 5 and 7, Fig. 2). This interaction of H37 with huCdc7 completely disappeared when antibody and peptide used as antigen for preparing the antibody were subjected to a pre-incubation in advance (cf. lane 8, Fig. 2). In extracts of HeLa cells, both anti-Cdc7 antibody and anti-H37 antibody were able to coprecipitate one polypeptide of 80 kDa that specifically reacted with anti-H37 antibody (lanes 9-13, Fig. 2).

[0028]

From the above result, it was found that intrinsic huCdc7 and H37 protein were present in cells as a complex.

Example 2

In order to investigate whether H37 protein has an ability of activating huCdc7, huCdc7/H37 complex prepared by expression of myc-tagged H37 and wild-type or kinase negative huCdc7 in Cos cells was immunoprecipitated by anti-huCdc7 antibody or anti-myc antibody and then a kinase reaction in vitro was measured using GST-MCM3 fusion protein as a substrate. The result is as shown in Fig. 3 and in Fig. 4. In the presence of wild-type huCdc7, phosphorylation in a good efficiency was observed for the MCM3 protein in both immunoprecipitate of anti-huCdc7 antibody and immunoprecipitate of anti-myc antibody (cf. lanes 2 and 7, Fig. 3). Further, two more phosphorylated proteins were observed and they were identified as transfected huCdc7 and myc-tagged H37 (data not shown). Since the phosphorylation as such was not detected at all in kinase negative huCdc7, it was confirmed

that kinase activity of huCdc7 acted in the phosphorylation. However, kinase negative huCdc7 was also able to form a complex with H37 protein (cf. lanes 3 and 8, Fig. 3; lanes 2 and 4, Fig. 4). In addition, the mobility of H37 protein on gel electrophoresis was slow when a wild-type huCdc7 was co-expressed and detected as plural bands while such a shift was not observed in kinase negative huCdc7 (cf. lanes 1 and 3, Fig. 4). The band where the mobility was slow disappeared by a treatment with phosphatase and, therefore, it was confirmed to be a perphosphorylated H37 protein (data not shown). In addition, when huCdc7 and H37 protein were co-expressed in insect cells, it was possible to reconstitute a very strong kinase activity which was able to efficiently phosphorylate MCM2 and MCM3 proteins (data not shown).

[0029]

The above result shows that H37 protein activates huCdc7 kinase and further that H37 protein itself is phosphorylated by huCdc7.

Further, under those experimental conditions, intrinsic H37 protein level was too low and in the case where only huCdc7 catalytic subunit was expressed and, therefore, the kinase activity was little (cf. lanes 4 and 9, Fig. 3). From those facts, it was confirmed that H37 protein encoded regulatory cell unit of huCdc7 and specifically activated its kinase activity.

Example 3

An amino acid sequence of the H37 protein (SEQ ID NO: 1) was analyzed. The result was that, as shown in Fig. 5 and Fig. 6, an amino acid sequence region having a homology of 33% with budding yeast Dbf4 was found. This conserved domain (H37 motif C) was also present in the gene similar to H37 identified in rat, fruit fly and fission yeast (cf. Fig. 6). Further, another amino acid sequence of H37 (H37 motif N) was conserved in the H37-related gene in rat and fruit fly. However, the H37 motif N was not conserved on the Dbf4 protein of a budding yeast (cf. Fig. 6).

[0030]

In order to determine the region on H37 protein essential for binding with huCdc7, a series of deletion derivatives of H37 at N- and C-terminals as shown in Fig. 7 was prepared, each of them was

expressed in yeast as a fusion protein with Gal4 activated domain and an interaction of each deletion derivative with huCdc7 was investigated by means of a two-hybrid assay. The result is as shown in Fig. 8. As a result of deletion at N-terminal, even a deletion of 255 amino acids at N-terminal did not affect the interaction with huCdc7 ($\Delta N2$). However, when 50 amino acids at N-terminal were further deleted whereupon the H37 motif C was deleted, an interaction with huCdc7 was completely lost ($\Delta N3$).

[0031]

On the other hand, with regard to deletion from C-terminal, when only 20 amino acids were deleted, the binding ability with huCdc7 decreased to an extent of about 60% (ΔC). In addition, when 243 or 369 amino acids were deleted from the C-terminal ($\Delta P2$ and ΔB), the interaction lowered to an extent of about 10% of the full-length clone. $\Delta P1$ containing only 235 amino acids of N-terminal did not interact with huCdc7. However, 50 amino acids that were commonly present in ΔB and $\Delta N2$ were not sufficient for an efficient interaction with huCdc7 (data not shown).

[0032]

A part of the above-mentioned H37 deletion derivatives was co-expressed in COS7 cells together with huCdc7 and it was confirmed by immunoprecipitation method whether it formed a complex with huCdc7 protein. The result was that, like in the result of a two-hybrid assay, it was confirmed that only H37 deletion derivatives of deltaB and deltaN2 formed a complex with huCdc7 (cf. Fig. 9).

[0033]

The above result shows that H37 motif-C is essential for an interaction of H37 protein with huCdc7 catalytic subunit but shows that H37 motif C alone is insufficient. In the budding yeast, it was reported already that the region containing H37 motif C was sufficient for binding with Cdc7 (Mol. Cell. Biol. 15:6775-6782, 1996). When an in vitro kinase reaction was carried out using those deletion derivatives, it was found that only 419 amino acids at the C-terminal containing Dbf4 motif-C was sufficient for activation of phosphorylating ability (data not shown).

Example 4

Expression patterns of H37 mRNA in various human tissues and cancer cells were investigated by northern blotting. The result is as shown in Fig. 10(a) and (b). By an H37 cDNA-specific probe, a transcribed product of 2.5 kb was detected in all tissues except brain and kidney and also in all cancer cells. That was contrary to the fact that, in brain and kidney, huCdc7 mRNA was relatively highly expressed (EMBO J. 16:4340-4351, 1997). Among the tissues tested, expression of H37 mRNA was highest in testicles and then in thymus, and it was reported by the inventors already that both of them were the tissues where the expression of huCdc7 catalytic subunit was particularly high as well (EMBO J. 16:4340-4351, 1997). In the testicles, two different RNA bands of 6 kb and 4 kb were also detected (cf. Fig. 10(a)) although it was ambiguous what the real substances therefor were. It was also confirmed that H37 mRNA was expressed in a very high level in nearly all cancer cells except lung cancer cell A549 (cf. Fig. 10(b)). That shows an important role of H37 protein in the cells having an active proliferating ability.

Example 5

In order to investigate whether expression of H37 was regulated by cell cycle, human normal fibroblast cell WI38 was synchronized with G0 phase by means of serum starvation (cf. Fig. 11), total RNA was prepared at various periods after addition of serum and the level of H37 mRNA was investigated by northern blotting. The result is as shown in Fig. 12. H37 mRNA level was low in the cell at the resting stage and, as the cells came near the boundary of G1 and S, it gradually increased and, after 20 hours from the serum addition, it became maximum. The expression pattern shown in Fig. 12 was similar to that of transcribed product of huCdc6 that is known to be induced by growth stimulation (Mol. Cell. Biol. 15:4215-4224, 1995; Proc. Natl. Acad. Sci. USA 94:142-147, 1997).

[0034]

Further, in order to investigate the changes in expression of H37 in cell cycle, human CEM cells were fractionated by means of elutriation method (cf. Fig. 13) and then a northern blotting was carried out (cf. Fig. 14). It was shown by the result thereof that H37 mRNA was low at G1 phase, increased from the late G1 phase to

S phase, highly maintained during S phase and somewhat decreased in G2 phase although it was still kept high. In an experiment where HeLa cells were stopped at G2 phase by nocodazole and the cell cycle was synchronously shifted (cf. Fig. 15), it was also found that H37 mRNA decreased together with shift from G2 to G1 and increased again at shift to S phase (cf. Fig. 16). Similarly, expression of Cdc6 rose together with shift from G1 phase to S phase but there was a difference from H37 in that, as S phase proceeded, it decreased and, at G2 phase it was suppressed to low. The result shows that the expression of H37 mRNA changes even in the proliferating cell cycle becomes maximum at S phase where it functions.

[0035]

In order to further investigate the expression of H37, intracellular localization of H37 protein in animal cells was measured. The result of indirect fluorescent antibody method using two kinds of H37-specific antibodies was that, in both HeLa and WI38 cells, intrinsic H37 protein was observed as very clear various spots in the nuclei (cf. Fig. 17).

When those results and the intranuclear localization of huCdc7 catalytic submit already reported by the inventors (EMBO J. 16:4340-4351, 1997) were taken into consideration, it was confirmed that huCdc7/H37 complex was a kinase localized in the nuclei and that its regulatory subunit H37 was expressed depending upon the stage of cell cycle.

Example 6

Function of the intrinsic H37 protein in the shift from G1 to S in cell cycle was investigated by antibody microinjection method. Antibody against N-terminal 305 amino acids of H37 protein (anti-H37N antibody) and antibody against C-terminal oligopeptide (anti-H37Cpep antibody) were treated by affinity purification and those antibodies were microinjected into normal fibroblast cells (KD cells) derived from human lip. KD cells were previously arrested at G0 phase by means of serum starvation and then synchronously proceeded in cell cycle by re-addition of serum. Numbers of the BrdU-positive cells, which BrdU is a nucleotide derivative and is incorporated into the cells, were measured whereby it was investigated how much fractions of cells

at S phase were present after various stages after addition of serum. The result is as shown in Fig. 18. It was confirmed that the cells started DNA synthesis after about 18 hours from serum addition and that, after 24 hours, almost 90% of the cells were in S phase.

5 [0036]

Accordingly, in this Example, antibody was microinjected at the period of after 12 hours after the addition of serum where the cells were still in G1 phase and, at 26 hours when cells would be of in S phase, the cells were fixed and the BrdU-positive cells were measured. The result is as shown in Fig. 19. Although 70% of the cells into which anti-H37N antibody was microinjected did not shift into S phase, there was nearly no effect by the control antibody. Even by the microinjection of the anti-H37Cpep antibody, an effect of inhibiting the shift to S phase which was same as or even more than the anti-H37N antibody was observed. In addition, when anti-H37Cpep antibody and a peptide which was an antigen for the preparation of anti-H37Cpep antibody were simultaneously microinjected into the cells, 70% or more of the cells were shifted to S phase.

[0037]

20 Fig. 20 is a staining example of BrdU and microinjected antibody. The antibody was microinjected at the stage when the cells were from the mid to late S phase and, during that period, expression of H37 protein is believed to be low. The microinjected antibody efficiently bound to the freshly synthesized H37 protein and, as a result, it is speculated to inhibit the transfer of H37 protein into nucleus. Those results strongly suggest the function of H37, i.e. the function of huCdc7/H37 complex is required for the progression of S phase of the animal cells.

[0038]

30 [Effects of the Invention]

As fully illustrated hereinabove, the present invention provided human H37 protein, a regulatory subunit for Cdc7 that regulate replication of human cells, human gene encoding this protein as well as cDNA thereof, an antibody against H37 protein, and a method for controlling the proliferation of human cells using those genetic engineering materials and antibody. As a result, it provides

potentially novel means for preparation of the necessary amount of stem cells or the like to be used for the therapy of various human diseases or for the suppression of proliferation of cancer cells.

[0039]

5 [Sequence Listing]

<110> Applicant Name: JAPAN SCIENCE AND TECHNOLOGY CORPORATION

<120> Title of Invention: HUMAN H37 PROTEIN AND cDNA ENCODING THE PROTEIN

<160> Number of Sequences: 4

10 <210> SEQ ID NO: 1

<211> Sequence Length: 674

<212> Sequence Type: PRT

<213> Homo sapiens

<400> Sequence Description

Met Asn Ser Gly Ala Met Arg Ile His Ser Lys Gly His Phe Gln Gly

1

5

10

15

Gly Ile Gln Val Lys Asn Glu Lys Asn Arg Pro Ser Leu Lys Ser Leu

15

	20		25		30
Lys Thr Asp Asn Arg Pro Glu Lys Ser Lys Cys Lys Pro Leu Trp Gly					
	35		40		45
Lys Val Phe Tyr Leu Asp Leu Pro Ser Val Thr Ile Ser Glu Lys Leu					
	50		55		60
Gln Lys Asp Ile Lys Asp Leu Gly Gly Arg Val Glu Glu Phe Leu Ser					
	65		70		75
Lys Asp Ile Ser Tyr Leu Ile Ser Asn Lys Lys Glu Ala Lys Phe Ala					
		85		90	
Gln Thr Leu Gly Arg Ile Ser Pro Val Pro Ser Pro Glu Ser Ala Tyr					
	100		105		110
Thr Ala Glu Thr Thr Ser Pro His Pro Ser His Asp Gly Ser Ser Phe					
	115		120		125
Lys Ser Pro Asp Thr Val Cys Leu Ser Arg Gly Lys Leu Leu Val Glu					
	130		135		140
Lys Ala Ile Lys Asp His Asp Phe Ile Pro Ser Asn Ser Ile Leu Ser					
	145		150		155
Asn Ala Leu Ser Trp Gly Val Lys Ile Leu His Ile Asp Asp Ile Arg					
		165		170	
Tyr Tyr Ile Glu Gln Lys Lys Lys Glu Leu Tyr Leu Leu Lys Lys Ser					
	180		185		190
Ser Thr Ser Val Arg Asp Gly Gly Lys Arg Val Gly Ser Gly Ala Gln					
	195		200		205
Lys Thr Arg Thr Gly Arg Leu Lys Lys Pro Phe Val Lys Val Glu Asp					
	210		215		220
Met Ser Gln Leu Tyr Arg Pro Phe Tyr Leu Gln Leu Thr Asn Met Pro					
	225		230		235
Phe Ile Asn Tyr Ser Ile Gln Lys Pro Cys Ser Pro Phe Asp Val Asp					
		245		250	
					255

Lys Pro Ser Ser Met Gln Lys Gln Thr Gln Val Lys Leu Arg Ile Gln
 260 265 270
 Thr Asp Gly Asp Lys Tyr Gly Gly Thr Ser Ile Gln Leu Gln Leu Lys
 275 280 285
 Glu Lys Lys Lys Lys Gly Tyr Cys Glu Cys Cys Leu Gln Lys Tyr Glu
 290 295 300
 Asp Leu Glu Thr His Leu Leu Ser Glu Gln His Arg Asn Phe Ala Gln
 305 310 315 320
 Ser Asn Gln Tyr Gln Val Val Asp Asp Ile Val Ser Lys Leu Val Phe
 325 330 335
 Asp Phe Val Glu Tyr Glu Lys Asp Thr Pro Lys Lys Lys Arg Ile Lys
 240 245 350
 Tyr Ser Val Gly Ser Leu Ser Pro Val Ser Ala Ser Val Leu Lys Lys
 355 360 365
 Thr Glu Gln Lys Glu Lys Val Glu Leu Gln His Ile Ser Gln Lys Asp
 370 375 380
 Cys Gln Glu Asp Asp Thr Thr Val Lys Glu Gln Asn Phe Leu Tyr Lys
 385 390 395 400
 Glu Thr Gln Glu Thr Glu Lys Lys Leu Leu Phe Ile Ser Glu Pro Ile
 405 410 415
 Pro His Pro Ser Asn Glu Leu Arg Gly Leu Asn Glu Lys Met Ser Asn
 420 425 430
 Lys Cys Ser Met Leu Ser Thr Ala Glu Asp Asp Ile Arg Gln Asn Phe
 435 440 445
 Thr Gln Leu Pro Leu His Lys Asn Lys Gln Glu Cys Ile Leu Asp Ile
 450 455 460
 Ser Glu His Thr Leu Ser Glu Asn Asp Leu Glu Glu Leu Arg Val Asp
 465 470 475 480
 His Tyr Lys Cys Asn Ile Gln Ala Ser Val His Val Ser Asp Phe Ser

	485	490	495
Thr Asp Asn Ser Gly Ser Gln Pro Lys Gln Lys Ser Asp Thr Val Leu			
	500	505	510
Phe Pro Ala Lys Asp Leu Lys Glu Lys Asp Leu His Ser Ile Phe Thr			
	515	520	525
His Asp Ser Gly Leu Ile Thr Ile Asn Ser Ser Gln Glu His Leu Thr			
	530	535	540
Val Gln Ala Lys Ala Pro Phe His Thr Pro Pro Glu Glu Pro Asn Glu			
545	550	555	560
Cys Asp Phe Lys Asn Met Asp Ser Leu Pro Ser Gly Lys Ile His Arg			
	565	570	575
Lys Val Lys Ile Ile Leu Gly Arg Asn Arg Lys Glu Asn Leu Glu Pro			
	580	585	590
Asn Ala Glu Phe Asp Lys Arg Thr Glu Phe Ile Thr Gln Glu Glu Asn			
	595	600	605
Arg Ile Cys Ser Ser Pro Val Gln Ser Leu Leu Asp Leu Phe Gln Thr			
	610	615	620
Ser Glu Glu Lys Ser Glu Phe Leu Gly Phe Thr Ser Tyr Thr Glu Lys			
625	630	635	640
Ser Gly Ile Cys Asn Val Leu Asp Ile Trp Glu Glu Glu Asn Ser Asp			
	645	650	655
Asn Leu Leu Thr Ala Phe Phe Ser Ser Pro Ser Thr Ser Thr Phe Thr			
	660	665	670
Gly Phe			
674			

<210> SEQ ID NO: 2

<211> Sequence Length: 234

<212> Sequence Type: PRT

<213> Homo sapiens

5 <400> Sequence Description

Met Asn Ser Gly Ala Met Arg Ile His Ser Lys Gly His Phe Gln Gly
 1 5 10 15
 Gly Ile Gln Val Lys Asn Glu Lys Asn Arg Pro Ser Leu Lys Ser Leu
 20 25 30
 Lys Thr Asp Asn Arg Pro Glu Lys Ser Lys Cys Lys Pro Leu Trp Gly
 35 40 45
 Lys Val Phe Tyr Leu Asp Leu Pro Ser Val Thr Ile Ser Glu Lys Leu
 50 55 60
 Gln Lys Asp Ile Lys Asp Leu Gly Gly Arg Val Glu Glu Phe Leu Ser
 65 70 75 80
 Lys Asp Ile Ser Tyr Leu Ile Ser Asn Lys Lys Glu Ala Lys Phe Ala
 85 90 95
 Gln Thr Leu Gly Arg Ile Ser Pro Val Pro Ser Pro Glu Ser Ala Tyr
 100 105 110
 Thr Ala Glu Thr Thr Ser Pro His Pro Ser His Asp Gly Ser Ser Phe
 115 120 125
 Lys Ser Pro Asp Thr Val Cys Leu Ser Arg Gly Lys Leu Leu Val Glu
 130 135 140
 Lys Ala Ile Lys Asp His Asp Phe Ile Pro Ser Asn Ser Ile Leu Ser
 145 150 155 160
 Asn Ala Leu Ser Trp Gly Val Lys Ile Leu His Ile Asp Asp Ile Arg
 165 170 175
 Tyr Tyr Ile Glu Gln Lys Lys Lys Glu Leu Tyr Leu Leu Lys Lys Ser
 180 185 190
 Ser Thr Ser Val Arg Asp Gly Gly Lys Arg Val Gly Ser Gly Ala Gln
 195 200 205
 Lys Thr Arg Thr Gly Arg Leu Lys Lys Pro Phe Val Lys Val Glu Asp
 210 215 220
 Met Ser Gln Ser Pro Ala Val His Leu Met
 225 230 234

<210> SEQ ID NO: 3

<211> Sequence Length: 2780

<212> Sequence Type: DNA

<213> Homo sapiens

5 <400> Sequence Description

AATTCGGCAC GAGCTCTCTG AGGCTGCGCC AAGACCTGAA GCGGCGGACC GAGAGCCCGG	60
GTCTGAGACT GAGAGAGCAA CGGAATGGAG GCGGGGTAGA GGCGGAAACA CAACCTGCAG	120
GGCCAGAGCG AGGCGCGAGA AGGACGGCGG CGTGAGGGGG CGGGGCGCGC AGCGCGAGAA	180
GGCAGGCACG AGGGGCGAGC GCGAGGCGGG GCACGGCGCG TGGCGTGAGA CGGGGCGGGG	240
CGCGCGTATC GGCGCCGCGG CCGCGTGACG CGTTTTCAAA TCTTCAACCG CCGCAGCCCA	300
CTCGTTTGTG CTTTGCGCCT TCCTCCTCCG CGCCTTGGAG CCGGATCCGG CCCCAGAAAC	360
CCGACCTGCA GACGCGGTAC CTCTACTGCG TAGAGGCCGT AGCTGGCGGA AGGAGAGAGG	420
CGGCCGTCCT GTCAACAGGC CGGGGGAAGC CGTGCTTTTCG CGGCTGCCCG GTGCGACACT	480
TTCTCCGGAC CCAGCATGTA GGTGCCGGGC GACTGCCATG AACTCCGGAG CCATGAGGAT	540
CCACAGTAAA GGACATTTCC AGGGTGGAAT CCAAGTCAAA AATGAAAAAA ACAGACCATC	600
TCTGAAATCT CTGAAAACTG ATAACAGGCC AGAAAAATCC AAATGTAAGC CACTTTGGGG	660
AAAAGTATTT TACCTTGACT TACCTTCTGT CACCATATCT GAAAAACTTC AAAAGGACAT	720
TAAGGATCTG GGAGGGCGAG TTGAAGAATT TCTCAGCAAA GATATCAGTT ATCTTATTTT	780
AAATAAGAAG GAAGCTAAAT TTGCACAAAC CTTGGGTCTGA ATTTCTCCTG TACCAAGTCC	840
AGAATCTGCA TATACTGCAG AAACCACTTC ACCTCATCCC AGCCATGATG GAAGTTCATT	900
TAAGTCACCA GACACAGTGT GTTTAAGCAG AGGAAAATTA TTAGTTGAAA AAGCTATCAA	960
GGACCATGAT TTTATTCCTT CAAATAGTAT ATTATCAAAT GCCTTGTCAT GGGGAGTAAA	1020
AATTC TTCAT ATTGATGACA TTAGATACTA CATTGAACAA AAGAAAAAAG AGTTGTATTT	1080
ACTCAAGAAA TCAAGTACTT CAGTAAGAGA TGGGGGCAAA AGAGTTGGTA GTGGTGCACA	1140
AAAAACAAGA ACAGGAAGAC TCAAAAAGCC TTTTGTAAG GTGGAAGATA TGAGCCAACT	1200
TTATAGGCCA TTTTATCTTC AGCTGACCAA TATGCCTTTT ATAAATTATT CTATTCAGAA	1260
GCCCTGCAGT CCATTTGATG TAGACAAGCC ATCTAGTATG CAAAAGCAAA CTCAGGTAA	1320

```

ACTAAGAATC CAAACAGATG GCGATAAGTA TGGTGAACC TCAATTCAAC TCCAGTTGAA 1380
AGAGAAGAAG AAAAAAGGAT ATTGTGAATG TTGCTTGCAG AAATATGAAG ATCTAGAAAC 1440
TCACCTTCTA AGTGAGCAAC ACAGAAACTT TGCACAGAGT AACCAGTATC AAGTTGTTGA 1500
TGATATTGTA TCTAAGTTAG TTTTGTACTT TGTGGAATAT GAAAAGGACA CACCTAAAAA 1560
GAAAAGAATA AAATACAGTG TTGGATCCCT TTCTCCTGTT TCTGCAAGTG TCCTGAAAAA 1620
GACTGAACAA AAGGAAAAAG TGAATTGCA ACATATTTCT CAGAAAGATT GCCAGGAAGA 1680
TGATACAACA GTGAAGGAGC AGAATTTCTT GTATAAAGAG ACCCAGGAAA CTGAAAAAAA 1740
GCTCCTGTTT ATTTTCAGAGC CCATCCCCCA CCCTTCAAAT GAATTGAGAG GGCTTAATGA 1800
GAAAATGAGT AATAAATGTT CCATGTTAAG TACAGCTGAA GATGACATAA GACAGAATTT 1860
TACACAGCTA CCTCTACATA AAAACAAACA GGAATGCATT CTTGACATTT CCGAACACAC 1920
ATTAAGTGAA AATGACTTAG AAGAATAAG GGTAGATCAC TATAAATGTA ACATACAGGC 1980
ATCTGTACAT GTTTCTGATT TCAGTACAGA TAATAGTGA TCTCAACCAA AACAGAAGTC 2040
AGATACTGTG CTTTTCCAG CAAAGGATCT CAAGGAAAAG GACCTTCATT CAATATTTAC 2100
TCATGATTCT GGTCTGATAA CAATAACAG TTCACAAGAG CACCTAACTG TTCAGGCAAA 2160
GGCTCCATTC CATACTCCTC CTGAGGAACC CAATGAATGT GACTTCAAGA ATATGGATAG 2220
TTTACCTTCT GGTAAAATAC ATCGAAAAGT GAAAATAATA TTAGGACGAA ATAGAAAAGA 2280
AAATCTGGAA CCAAATGCTG AATTTGATAA AAGAACTGAA TTTATTACAC AAGAAGAAAA 2340
CAGAATTTGT AGTTCACCGG TACAGTCTTT ACTAGACTTG TTTCAGACTA GTGAAGAGAA 2400
ATCAGAATTT TTGGGTTTCA CAAGCTACAC AGAAAAGAGT GGTATATGCA ATGTTTTAGA 2460
TATTTGGGAA GAGGAAAATT CAGATAATCT GTTAACAGCG TTTTCTCGT CCCCTTCAAC 2520
TTCTACATTT ACTGGCTTTT AGAATTTAAA AAATGCATAC TTTTCAGAAG TGATAAGGAT 2580
CATATTCCTG AAATTTTTAT AAATATGTAT GGAAATTCCT AGGATTTTTT TACCAGCTTT 2640
GTTTACAGAC CCAAATGTAA ATATTAAAAA TAAATATTTG CAATTTTCTA CAGAATTGAA 2700
TACCTGTTAA AGAAAAATTA CAGAATAAAC TTGTGACTGG TCTTGTTTTA CATTAAAAAA 2760
AAAAAAAAAA AAAACTCGAG 2780

```

<210> SEQ ID NO: 4

<211> Sequence Length: 2719

<212> Sequence Type: DNA

5 <213> Homo sapiens

<400> Sequence Description

AATTCGGCAC GAGCTCTCTG AGGCTGCGCC AAGACCTGAA GCGGCGGACC GAGAGCCCGG	60
GTCTGAGACT GAGAGAGCAA CGGAATGGAG GCGGGGTAGA GGCGGAAACA CAACCTGCAG	120
GGCCAGAGCG AGGCGCGAGA AGGACGGCGG CGTGAGGGGG GGGGCGCGC AGCGCGAGAA	180
GGCAGGCACG AGGGGCGAGC GCGAGGCGGG GCACGGCGCG TGGCGTGAGA CGGGGCGGGG	240
CGCGCGTATC GGCGCCGCGG CCGCGTGACG CGTTTTCAAA TCTTCAACCG CCGCAGCCCA	300
CTCGTTTGTG CTTTGCGCCT TCCTCCTCCG CGCCTTGGAG CCGGATCCGG CCCCCGAAAC	360
CCGACCTGCA GACGCGGTAC CTCTACTGCG TAGAGGCCGT AGCTGGCGGA AGGAGAGAGG	420
CGGCCGTCCT GTCAACAGGC CGGGGGAAGC CGTGCTTTTCG CGGCTGCCCG GTGCGACACT	480
TTCTCCGGAC CCAGCATGTA GGTGCCGGGC GACTGCCATG AACTCCGGAG CCATGAGGAT	540
CCACAGTAAA GGACATTTCC AGGGTGGAAT CCAAGTCAAA AATGAAAAAA ACAGACCATC	600
TCTGAAATCT CTGAAAAC TG ATAACAGGCC AGAAAAATCC AAATGTAAGC CACTTTGGGG	660
AAAAGTATTT TACCTTGA CT TACCTTCTGT CACCATATCT GAAAAACTTC AAAAGGACAT	720
TAAGGATCTG GGAGGGCGAG TTGAAGAATT TCTCAGCAAA GATATCAGTT ATCTTATTTT	780
AAATAAGAAG GAAGCTAAAT TTGCACAAAC CTTGGGTCTGA ATTTCTCCTG TACCAAGTCC	840
AGAATCTGCA TATACTGCAG AAACCACTTC ACCTCATCCC AGCCATGATG GAAGTTCATT	900
TAAGTCACCA GACACAGTGT GTTTAAGCAG AGGAAAATTA TTAGTTGAAA AAGCTATCAA	960
GGACCATGAT TTTATTCCTT CAAATAGTAT ATTATCAAAT GCCTTGTCAT GGGGAGTAAA	1020
AATTCTTCAT ATTGATGACA TTAGATACTA CATTGAACAA AAGAAAAAAG AGTTGTATTT	1080
ACTCAAGAAA TCAAGTACTT CAGTAAGAGA TGGGGGCAAA AGAGTTGGTA GTGGTGACAA	1140
AAAAACAAGA ACAGGAAGAC TCAAAAAGCC TTTTGTAAG GTGGAAGATA TGAGCCAAAG	1200
CCCTGCAGTC CATTTGATGT AGACAAGCCA TCTAGTATGC AAAAGCAAAC TCAGGTAAAA	1260
CTAAGAATCC AAACAGATGG CGATAAGTAT GGTGGAACCT CAATTCAACT CCAGTTGAAA	1320
GAGAAGAAGA AAAAAGGATA TTGTGAATGT TGCTTGACAGA AATATGAAGA TCTAGAAACT	1380
CACCTTCTAA GTGAGCAACA CAGAACTTT GCACAGAGTA ACCAGTATCA AGTTGTTGAT	1440
GATATTGTAT CTAAGTTAGT TTTTGACTTT GTGGAATATG AAAAGGACAC ACCTAAAAAG	1500
AAAAGAATAA AATACAGTGT TGGATCCCTT TCTCCTGTTT CTGCAAGTGT CCTGAAAAAG	1560
ACTGAACAAA AGGAAAAAGT GGAATTGCAA CATATTTCTC AGAAAGATTG CCAGGAAGAT	1620
GATACAACAG TGAAGGAGCA GAATTTCTCTG TATAAAGAGA CCCAGGAAAC TGAAAAAAG	1680

```

CTCCTGTTTA TTTCAGAGCC CATCCCCAC CCTTCAAATG AATTGAGAGG GCTTAATGAG 1740
AAAATGAGTA ATAAATGTTC CATGTTAAGT ACAGCTGAAG ATGACATAAG ACAGAATTTT 1800
ACACAGCTAC CTCTACATAA AAACAAACAG GAATGCATTC TTGACATTTT CGAACACACA 1860
TTAAGTGAAA ATGACTTAGA AGAACTAAGG GTAGATCACT ATAAATGTAA CATAACAGGCA 1920
TCTGTACATG TTTCTGATTT CAGTACAGAT AATAGTGGAT CTCAACCAAA ACAGAAGTCA 1980
GATACTGTGC TTTTTCAGC AAAGGATCTC AAGGAAAAGG ACCTTCATTC AATATTTACT 2040
CATGATTCTG GTCTGATAAC AATAAACAGT TCACAAGAGC ACCTAACTGT TCAGGCAAAG 2100
GCTCCATTCC ATACTCCTCC TGAGGAACCC AATGAATGTG ACTTCAAGAA TATGGATAGT 2160
TTACCTTCTG GTAAAATACA TCGAAAAGTG AAAATAATAT TAGGACGAAA TAGAAAAGAA 2220
AATCTGGAAC CAAATGCTGA ATTTGATAAA AGAACTGAAT TTATTACACA AGAAGAAAAC 2280
AGAATTTGTA GTTCACCGGT ACAGTCTTTA CTAGACTTGT TTCAGACTAG TGAAGAGAAA 2340
TCAGAATTTT TGGGTTTCAC AAGCTACACA GAAAAGAGTG GTATATGCAA TGTTTTAGAT 2400
ATTTGGGAAG AGGAAAATTC AGATAATCTG TTAACAGCGT TTTTCTCGTC CCCTTCAACT 2460
TCTACATTTA CTGGCTTTTA GAATTTAAAA AATGCATACT TTTCAGAAAGT GATAAGGATC 2520
ATATTCTTGA AATTTTATA AATATGTATG GAAATTCTTA GGATTTTTTT ACCAGCTTTG 2580
TTTACAGACC CAAATGTAAA TATTAATAAT AAATATTTGC AATTTTCTAC AGAATTGAAT 2640
ACCTGTAAAA GAAAAATTAC AGAATAAACT TGTGACTGGT CTTGTTTTAC ATTAAAAAAA 2700
AAAAAATAAA AACTCGAG 2719

```

[Brief Description of the Drawings]

[Figure 1]

5 This is the result of a western blotting measuring the
coimmunoprecipitation of H37 with huCdc7 expressed in mammalian cells.
Lanes 1-4, immunoprecipitates; lanes 5-7, whole cell extract. Upper
and middle panels, immunoprecipitated with anti-huCdc7 antibody No.
1; lower panel, immunoprecipitated with anti-myc antibody. Extracts
10 were prepared from Cos7 cells transfected with huCdc7 in combination
with H 1 (lanes 2 and 5), H 18 (lanes 3 and 6), H37 (lanes 4 and 7)
or huCdc7 alone (lane 1). Western blotting was conducted with
anti-myc antibody (upper panel) or an anti-huCdc7 antibody No. 1
(middle and lower panel).

15 [Figure 2]

This is a result of the western blotting measuring antibodies against H37 protein and association of huCdc7 and H37 *in vivo*. Nuclear extracts prepared from Cos7 cells transfected with myc-tagged H37 were blotted with anti-H37C (lane 1), anti-H37N (lane 2), anti-H37Cpep (lane 3) or anti-myc (lane 4) antibody. The arrow indicates the myc-tagged H37 protein, which carries 63 amino acids derived from 5' non-coding region in addition to the myc-tag. Immunoprecipitations from CEM extracts using either anti-huCdc7Cpep (lanes 5 and 6) or anti-H37Cpep (lanes 7 and 8) were separated on gel electrophoresis and blotted with huCdc7 monoclonal antibody (4A8). The symbols - and + indicates absence and presence, respectively, of each antigen peptide during immunoprecipitation. Lanes 9-13; immunoprecipitates, prepared from nuclear extracts of HeLa cells by anti-huCdc7 No. 1 (lane 9), anti-huCdc7 monoclonal antibody 4A8 (lane 10), anti-H37C (lane 11), anti-H37N (lane 12) or anti-H37Cpep (lane 13), were blotted with anti-H37Cpep.

[Figure 3]

This shows the result of immunoprecipitation by the use of anti-huCdc7 antibody No. 1 (lanes 1-5) or anti-myc antibody (lanes 6-10) from the extract of Cos7 cells transfected with myc-tagged H37 alone (lanes 1 and 6), or together with wild-type huCdc7 (lanes 2 and 7), or kinase negative huCdc7 (lanes 3 and 8). Wild-type huCdc7 alone (lanes 4 and 9) and kinase negative huCdc7 alone (lanes 5 and 10) were included as control.

[Figure 4]

This shows mobility shift of H37 induced by coexpression of wild-type huCdc7. Extracts were prepared from Cos7 cells expressing either wild-type or kinase negative huCdc7 together with myc-tagged H37. Immunoprecipitates with anti-huCdc7 antibody No. 1 (lanes 1 and 2) or anti-myc antibody (lanes 3 and 4) were blotted with anti-myc antibody (upper) or anti-huCdc7 antibody (lower). In all cases, samples were run on 8% SDS-PAGE.

[Figure 5]

This shows an amino acid sequence of the full-length H37 protein which is as same as the SEQ ID NO: 1.

[Figure 6]

This shows schematic representation of conserved regions between Dbf4 and H37. The double-arrowed region on Dbf4 was reported to be sufficient for interaction with huCdc7. Solid and gray double-arrowed region on H37 indicates the portion essential for interaction with huCdc7 or that sufficient for activation of huCdc7 kinase activity, respectively.

[Figure 7]

This is a schematic representation of N-terminal and C-terminal deletion derivatives of H37 proteins. The number at the end of each bar indicates the portion of the amino acid (corresponding to SEQ ID NO: 1) at the deletion endpoint. The striped region indicates Dbf4 motif-C.

[Figure 8]

This shows lacZ activity of H37 deletion derivatives in two-hybrid assay with huCdc7.

[Figure 9]

This shows the result of antibody coprecipitation method in which H37 deletion derivatives and huCdc7 were co-expressed in COS7 cells and a complex formation was measured.

[Figure 10]

(a) is the result of northern analysis of H37 mRNA expression in various tissues, and (b) is the result of a northern analysis of H37 mRNA expression in various cancer cell lines.

[Figure 11]

This shows the results where WI38 cells in a resting phase are stimulated by addition of 10% of serum and DNA contents at various stages were analyzed by means of FACS.

[Figure 12]

This shows the result of northern analysis of H37 and huCdc6 expression by using RNA extracted from cells of Fig. 11 (upper), and the graphs showing the relative expression amount of each mRNA (middle and lower).

[Figure 13]

This shows graphs showing each cell cycle fraction of human CEM cells fractionated by an elutriation method.

[Figure 14]

This shows the result of northern analysis for expression of H37 and cyclin E in each fraction of Fig. 13 (upper), and also shows the graphs showing the relative expression amounts of each mRNA (middle and lower).

5 [Figure 15]

This shows the result of FACS analysis for DNA content in which HeLa cells were stopped at latter G2 phase using nocodazole and then the cell cycle was synchronously shifted.

[Figure 16]

10 This shows the result northern analysis for expression of H37 and cyclin E in each cell cycle of Fig. 14 (upper), and also shows the graphs showing the relative expression amounts of each mRNA (middle and lower).

[Figure 17]

15 This shows H37 localization in cells measured by indirect immunofluorescent technique. The antibodies used are anti-H37C (A), anti-H37N (C) and a control antibody (E) while B, D and F were pictures stained with DAPI.

[Figure 18]

20 This shows kinetics of induction of DNA replication after serum stimulation of KD cells as measured by BrdU incorporation.

[Figure 19]

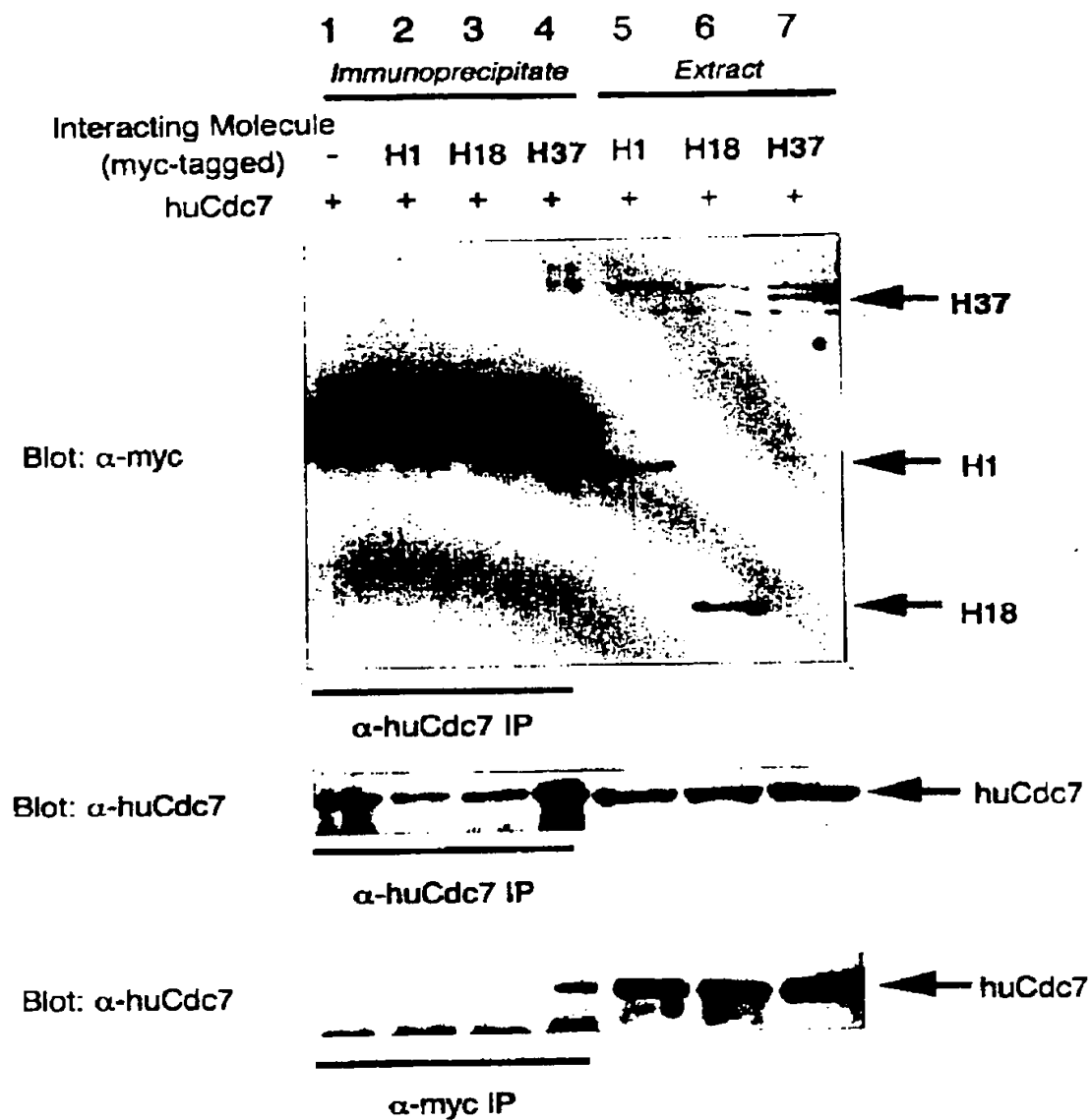
25 This shows the results in which KD cells, synchronized by serum starvation, were microinjected with various antibodies at 12 hours after serum stimulation, and then the rate of the cells incorporating the BrdU after 16 hours was measured. Thus, the rate of the cells which are conducting the DNA synthesis are shown.

[Figure 20]

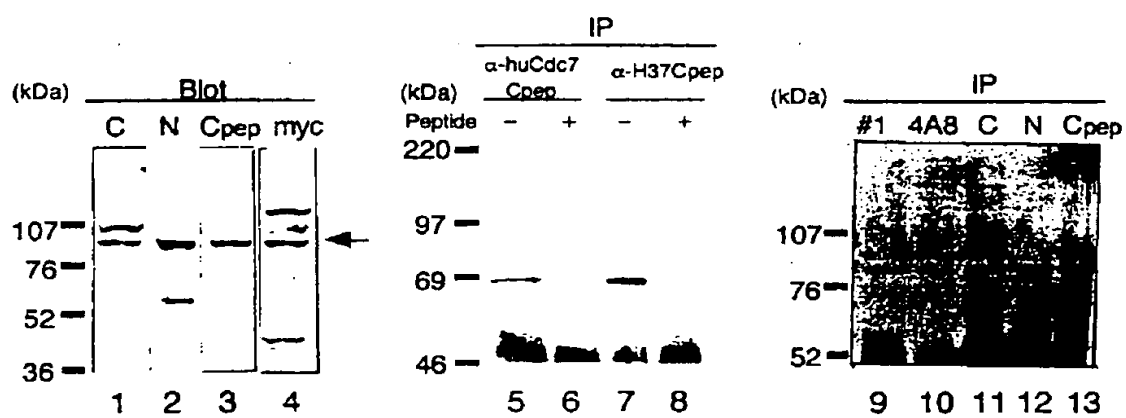
30 This is microscopic pictures of examples of cells into which a mixture of anti-H37Cpep antibody (left) or anti-H37Cpep and peptide was microinjected. The pictures show the incorporated BrdU (upper), injected antibody (middle) and cells (lower).

[Document Name] Drawings

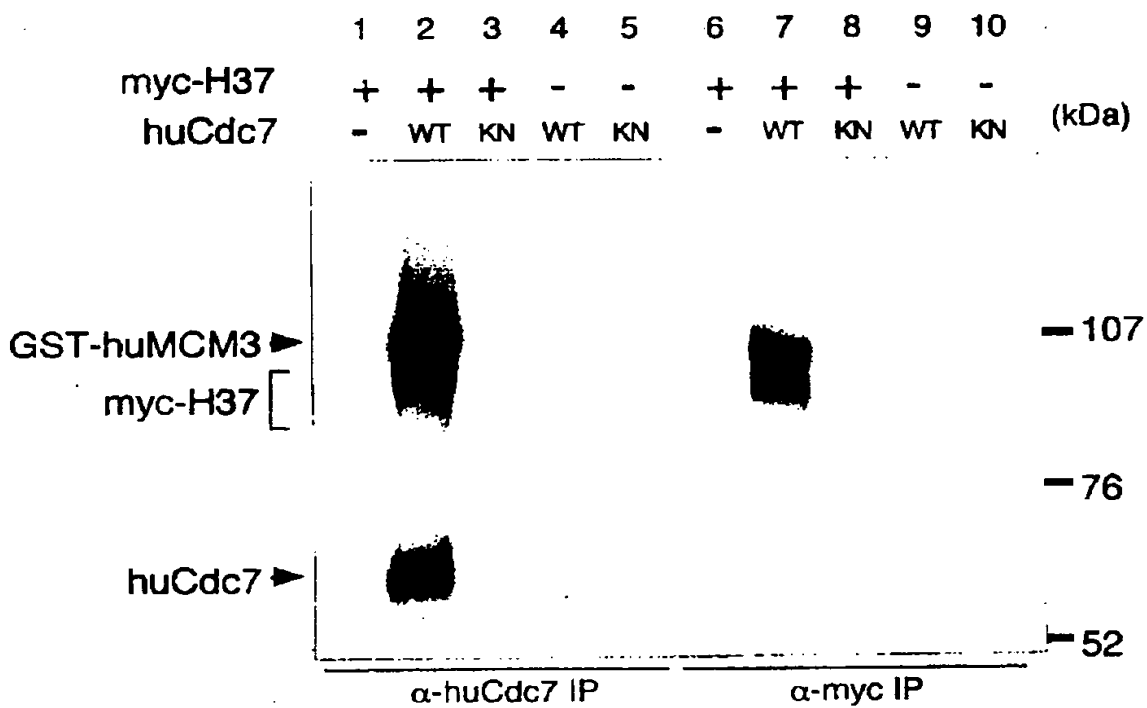
[Figure 1]



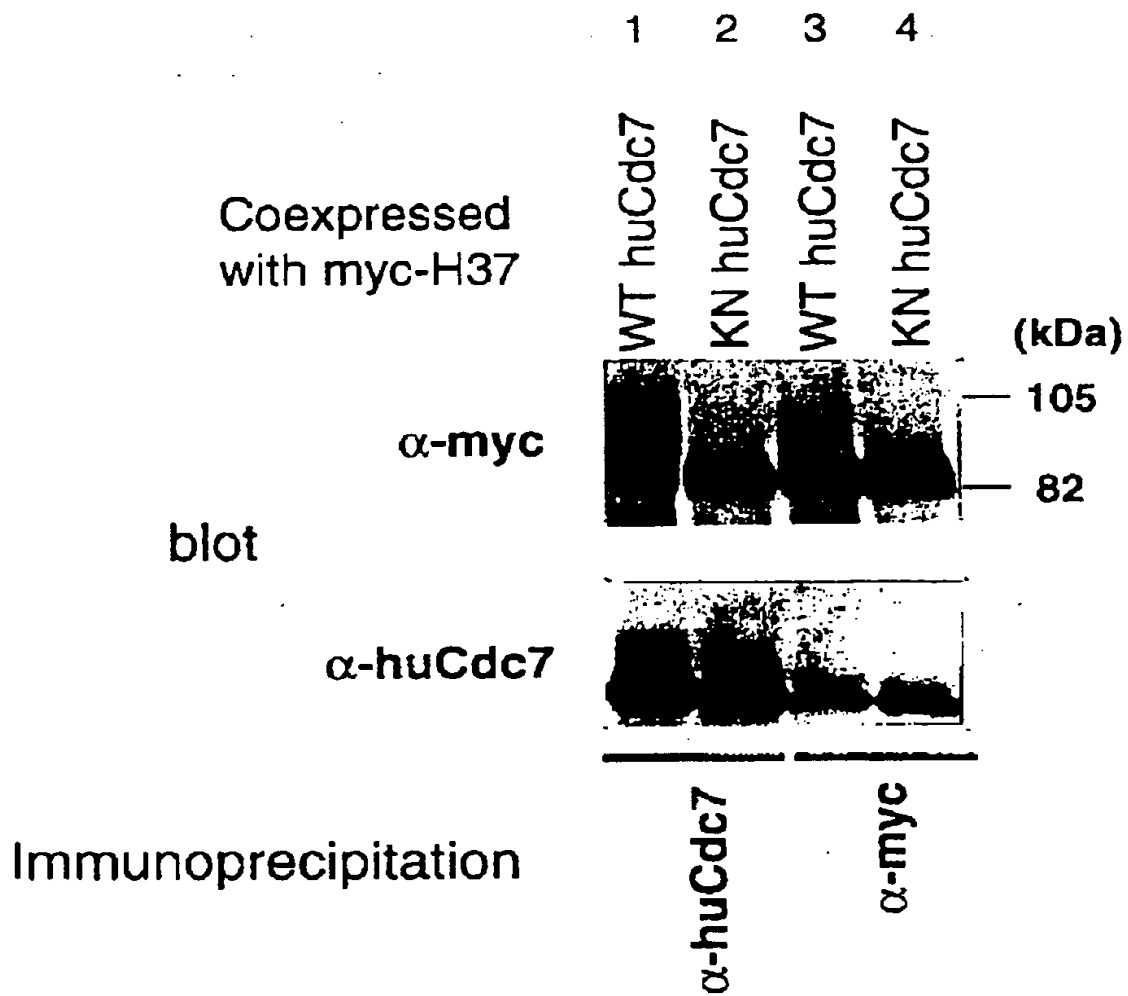
[Figure 2]



[Figure 3]



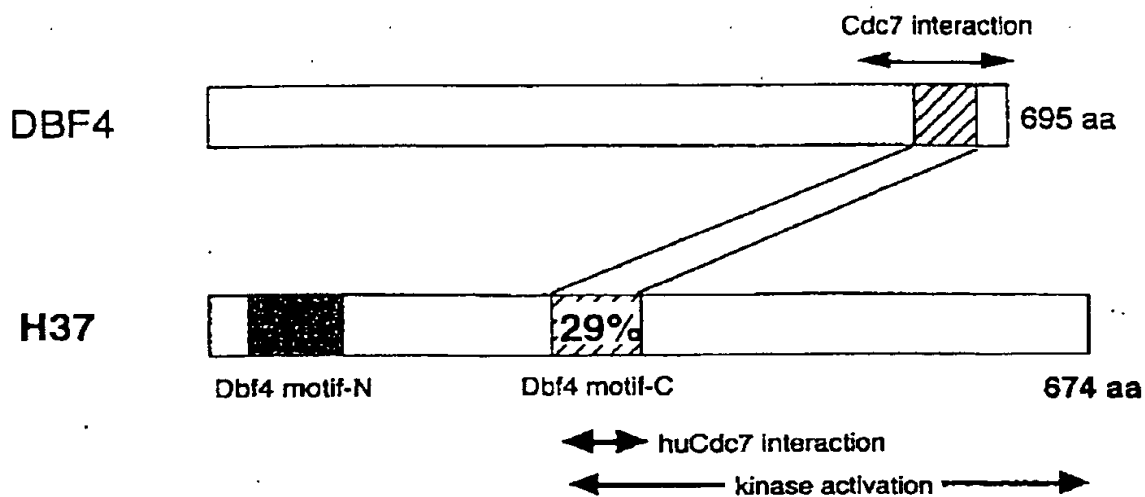
[Figure 4]



[Figure 5]

1 10 20 30 40 50 60
MMSGAMRIHSGHFGGGIQQVKNENRPSLKSLKTDNRPEKSKCKPLWGKVIFYLDLPSVTI
Dbf4 motif-N
61 70 80 90 100 110 120
SEKLOKDIKDLGGRVVEFLSKDISYLLSNKKEAKFAOTLGRISPVSPESAYTAETTSPH
121 130 140 150 160 170 180
PSHDGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPNSNLSNALSWGKILHIDDIRYYIE
181 190 200 210 220 230 240
QKKKELYLLKKSSTSVRDGGKRVGSGAQKTRTGRLLKPPVKVEDMSQLYRPFYLLQLTNP
241 250 260 270 280 290 300
FINYSIQKPCSPFDVDPSSMQKOTQVKLRIOTDGDKYGGTSIQLOLKEKKKGYCECCL
Dbf4 motif-C
301 310 320 330 340 350 360
QKYEDLETHLLSEQHRNFAQSNQYQVVDDIVSKLVDFVVEYEKDTPKKRKRIKYSVGSLSP
361 370 380 390 400 410 420
VSASVLKTEQKEKVELQHISQKDCQEDDTTVKEQNFLYKETQETEKLLFISEPIPHPS
421 430 440 450 460 470 480
NELRGLNEKMSNKCSMLSTAEDDIRQNFTQLPLKKNQECILDISEHTLSENDLEELRVD
481 490 500 510 520 530 540
HYKCNIQASVHVSDFDSTNDSGSPKQKSDTVLFPAKDLKEKDLHSIFTHDSGLITINSSQ
541 550 560 570 580 590 600
EHLTVQAKAPFHTPPPEPNEDCFKNMDSLPSGKIHRKVKIILGRNRKENLEPNAEFDKRT
601 610 620 630 640 650 660
EPITQEENRICSSPVQSLDLFQTSEEKSEFLGFTSYTEKSGICNVLDIWEENESDNLLT
661 670 674
AFFSSPSTSTFTGF*

[Figure 6]



Dbf4 motif-N

H37 32: **LTIRPRLS** **QEDWGV** **CVYIDDSV** **ETSE** **LDQDID** **GGGAL** **CLGSEISS** **LTGR** 91
 mu-H37 : **LTIRPRLS** **QEDWGV** **CVYIDDSV** **ETSE** **LDQDID** **GGGAL** **CLGSEISS** **LTGR**
 Dm-H37 : **TPPKVKV** **IKSE** **RP** **CHFK** **YER** **ICDHQLAKRIES** **DHALG** **HU** **FLL** **DDITH** **VTLP**

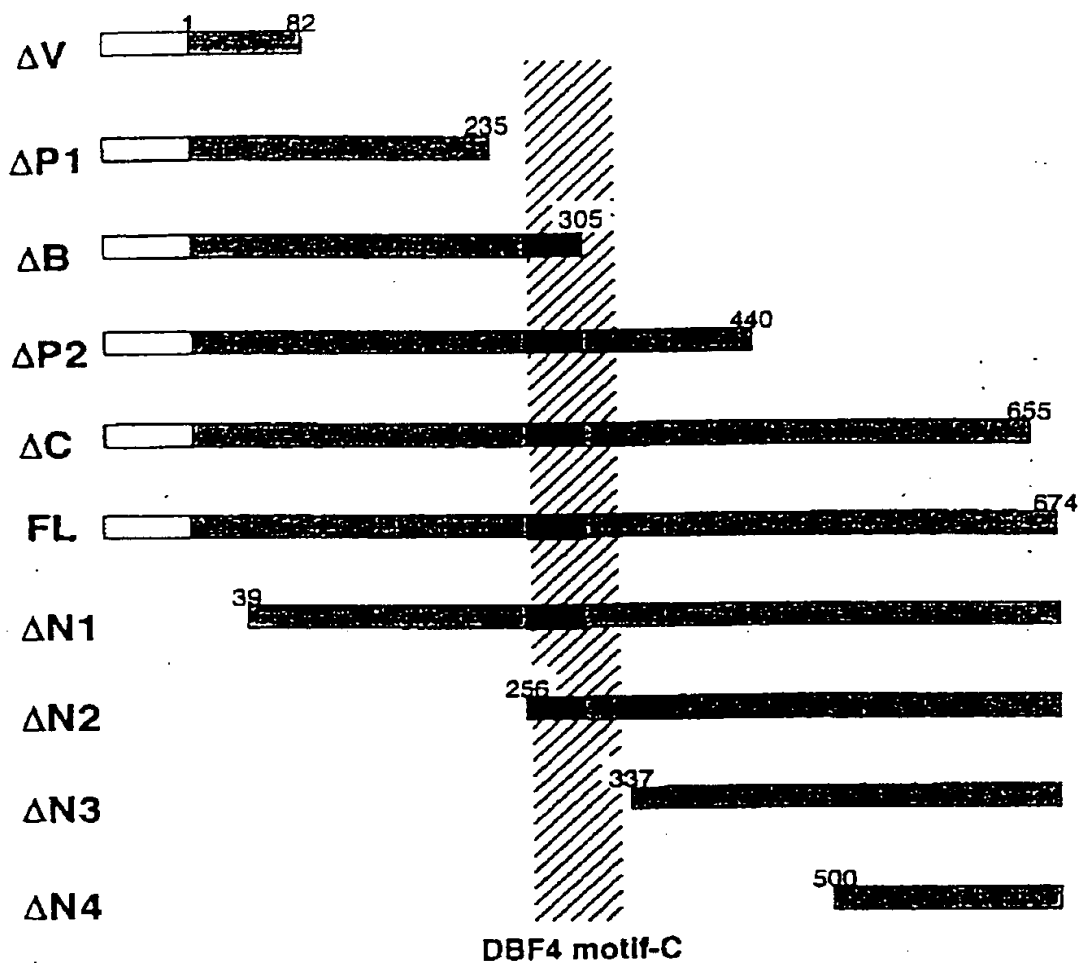
huDbf4N 92: **FAVFAQT** 98
 muDbf4N : **FAVFAQT**
 DmDbf4N : **EVIGGTS**

Dbf4 motif-C

H37 263: **QTCV** **KLRIQ** **TDGDKY** **GT** **SIQLQLKEK** **KKG** **YCF** **CLQRY** **DLIT** **TL** **STC** **RNF** **AQSS** 322
 mu-H37 : **QTCV** **KLRIQ** **TDGDKC** **GT** **SIQLQLKEK** **KKG** **YCF** **CLQRY** **DLIT** **TL** **STC** **RNF** **AQSS**
 Dm-H37 : **PSL** **DELKKQ** **SAIPNS** **PRSN** **CREPID** **SSER** **QCCV** **CL** **KLEYD** **IN** **HT** **OSK** **DEL** **AKNS**
 Dbf4 619: **IKST** **STNVT** **LHFNAQT** **AC** **IAQPV** **KKET** **ANS** **YCH** **NRVE** **YS** **CH** **IV** **SL** **HL** **SPAE** 677

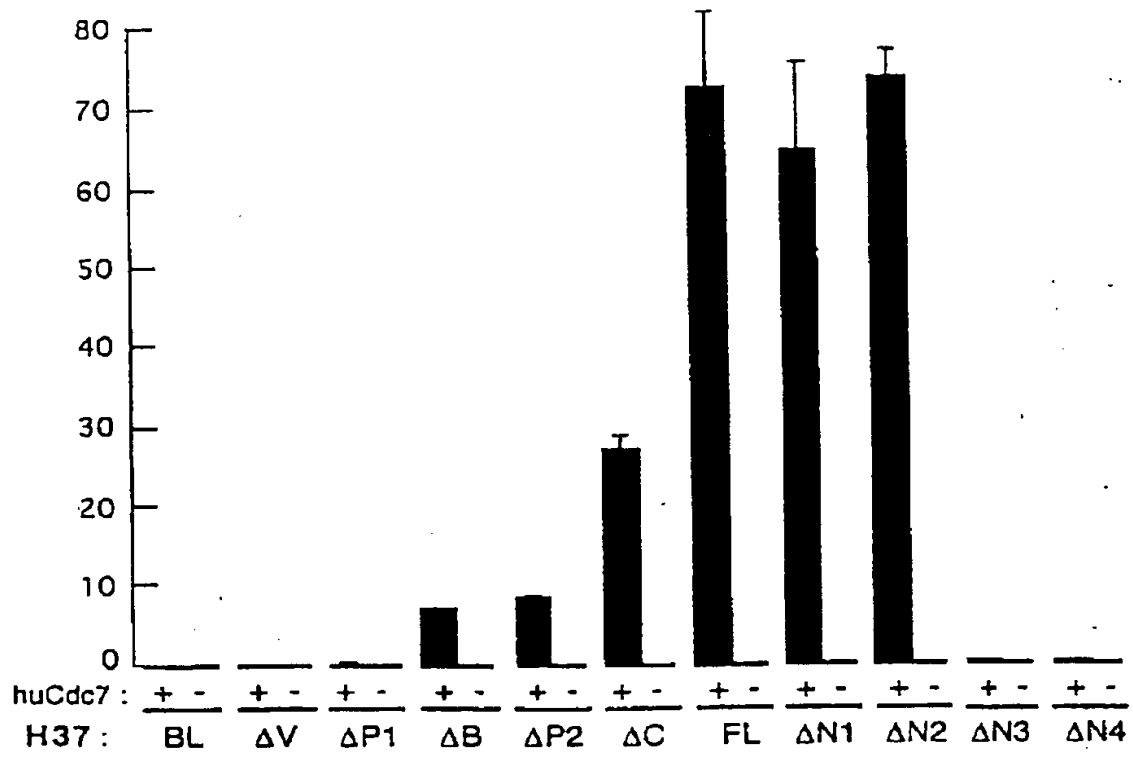
[Figure 7]

GAL4-AD

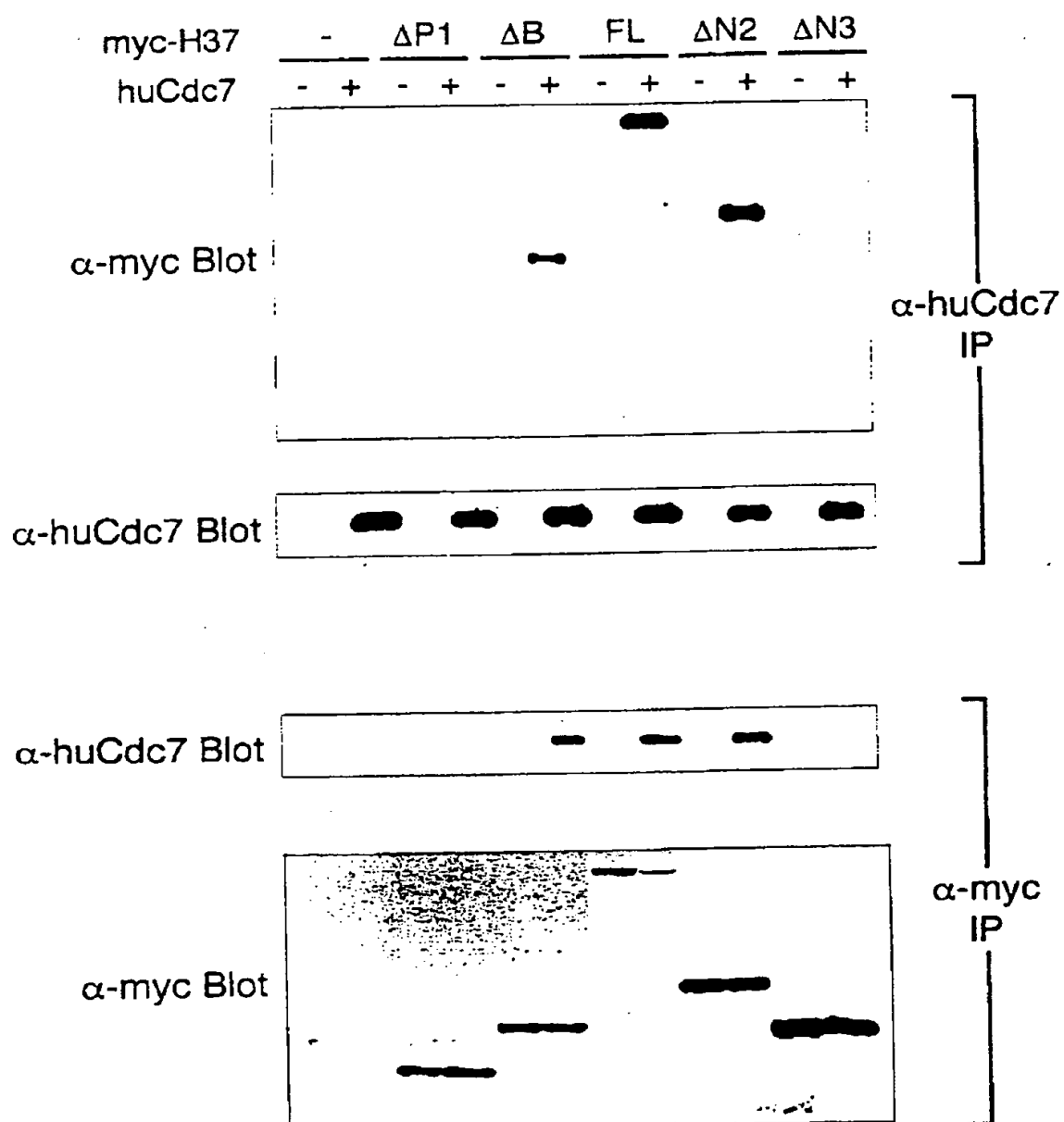


[Figure 8]

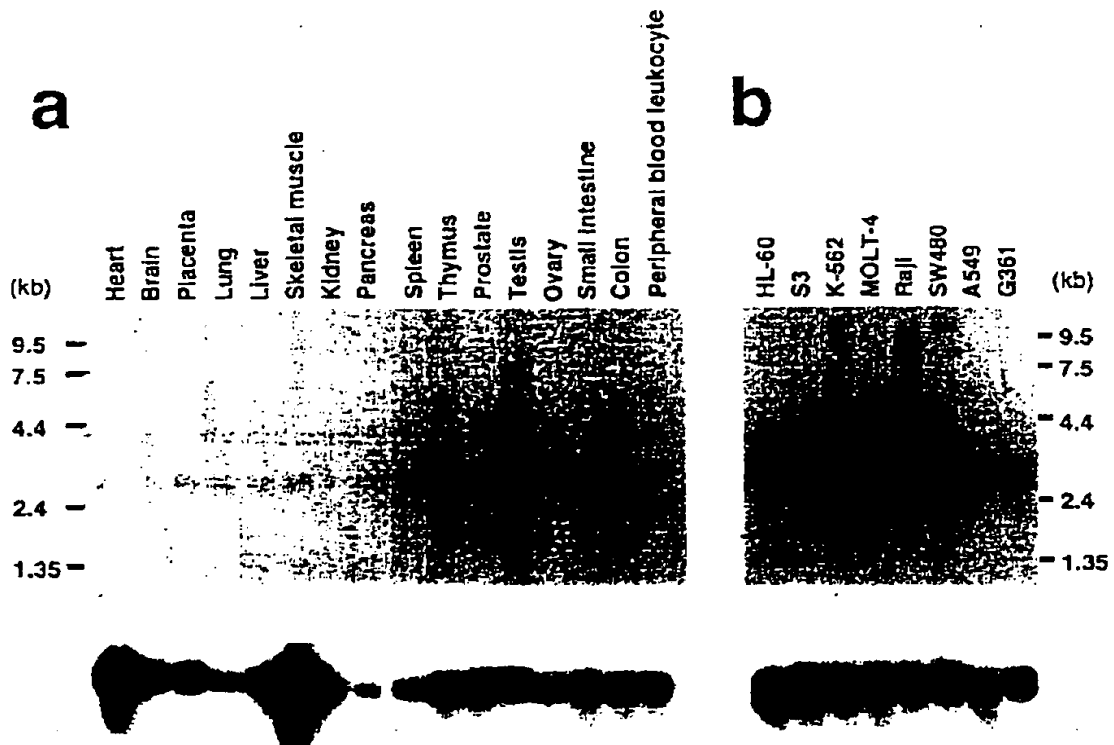
β -galactosidase
Activity (Unit)



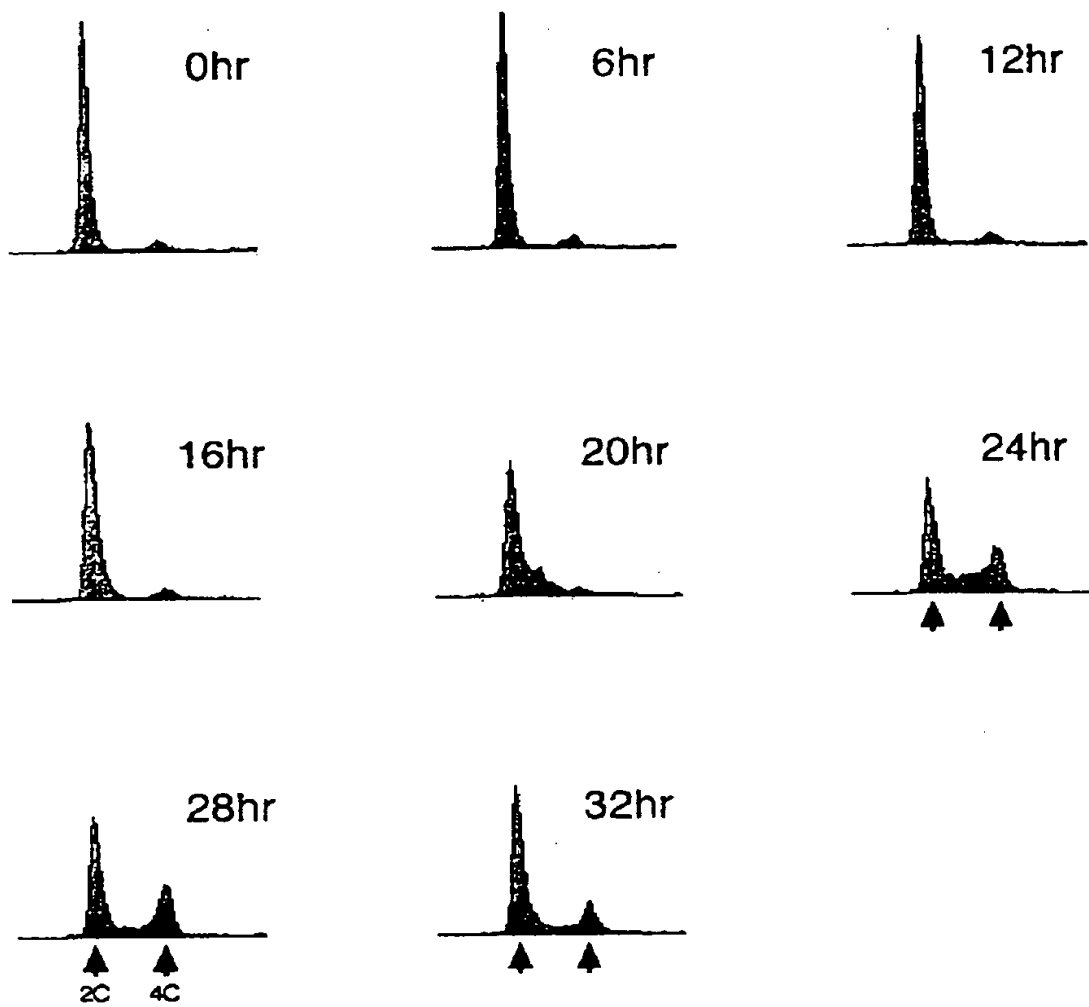
[Figure 9]



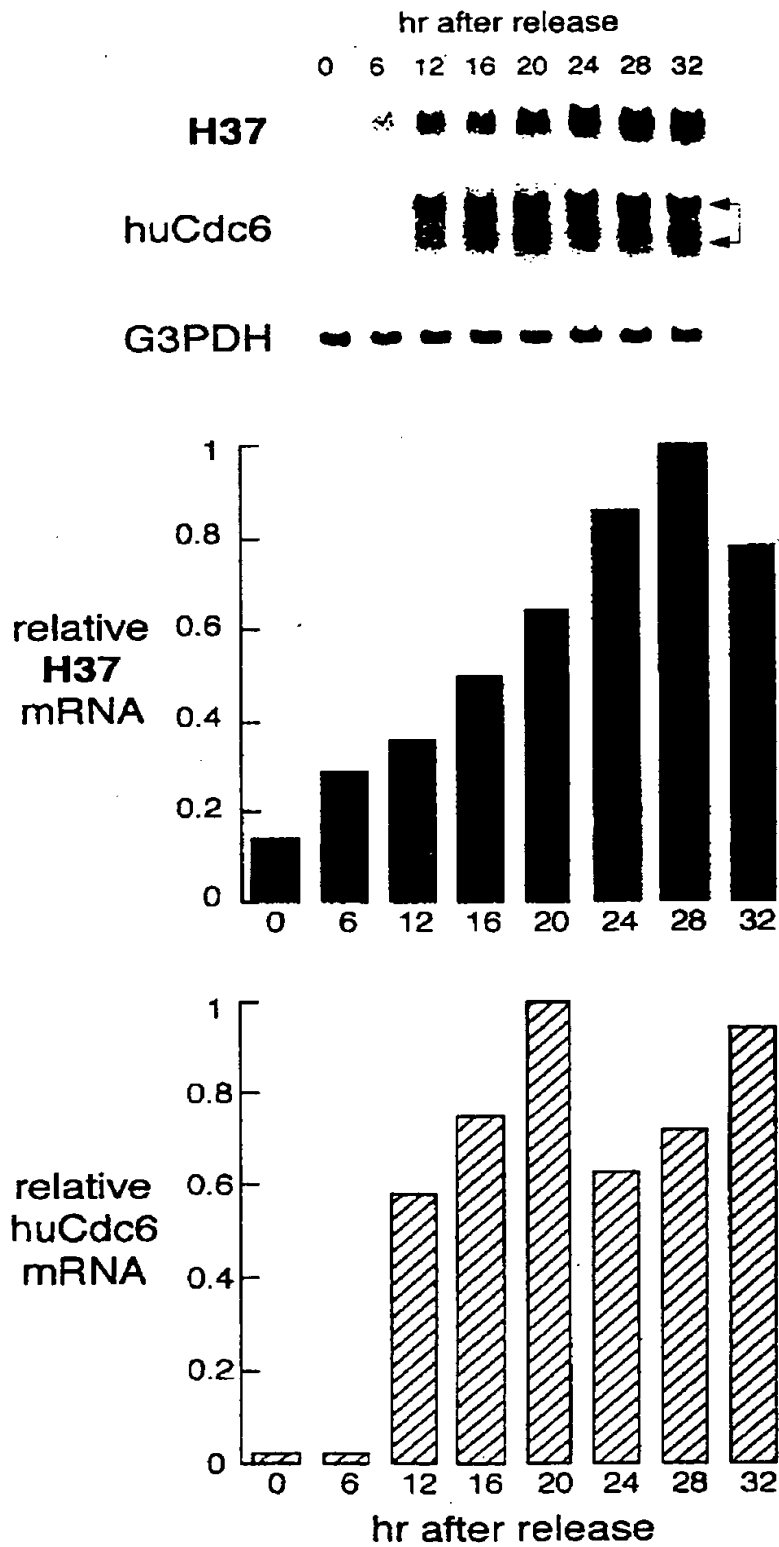
[Figure 10]



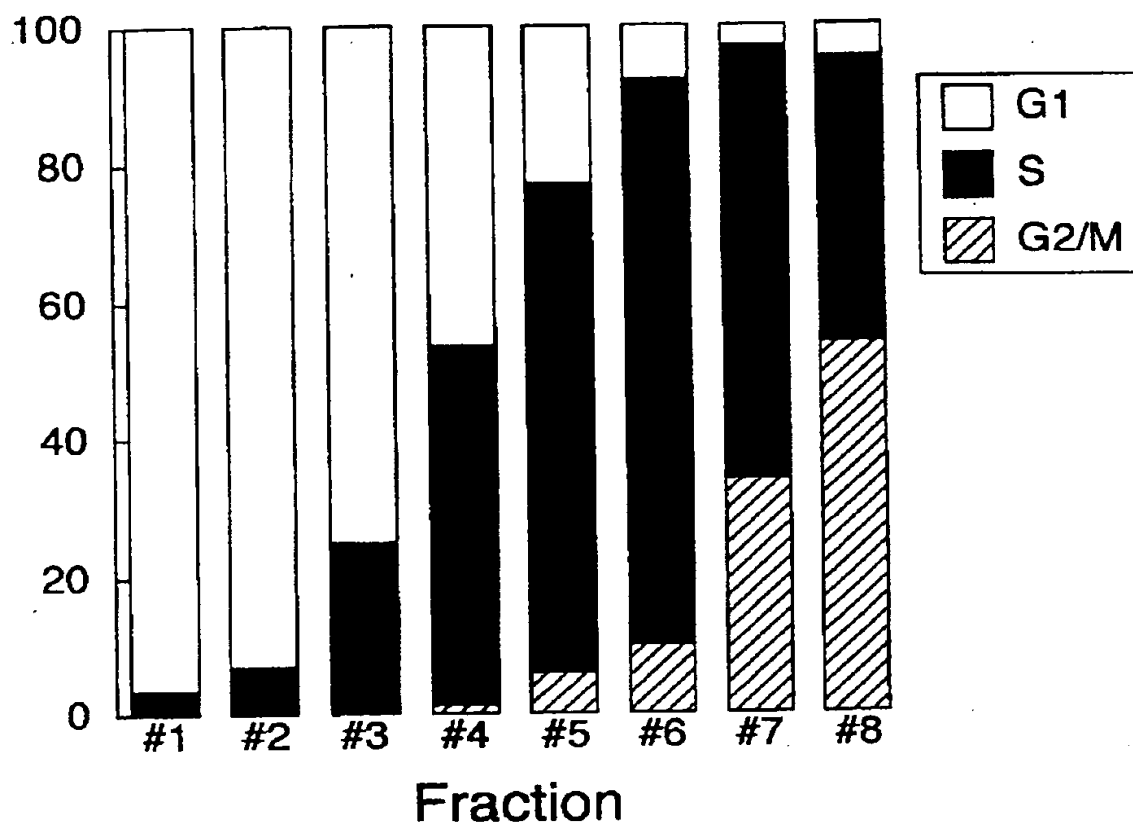
[Figure 11]



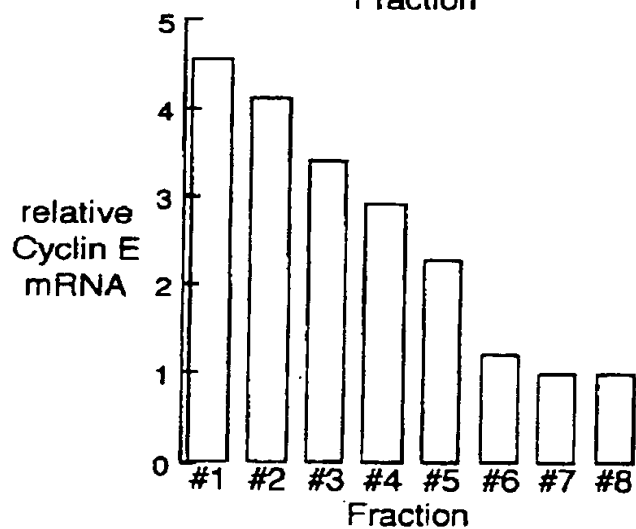
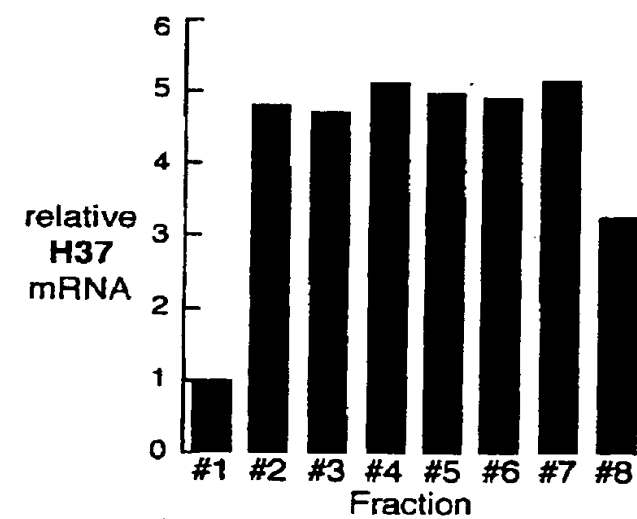
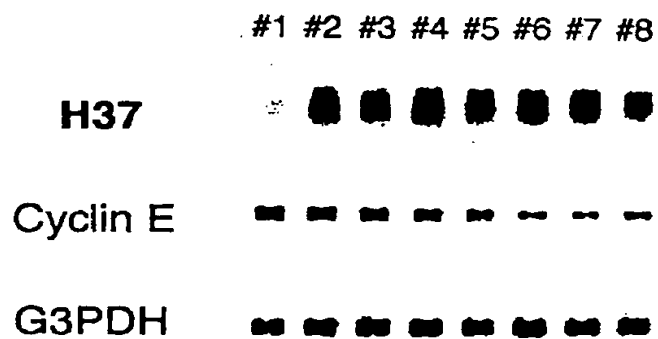
[Figure 12]



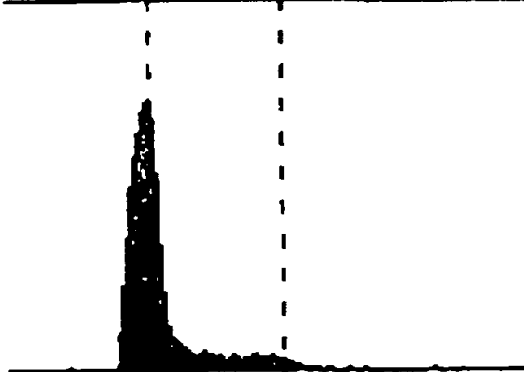
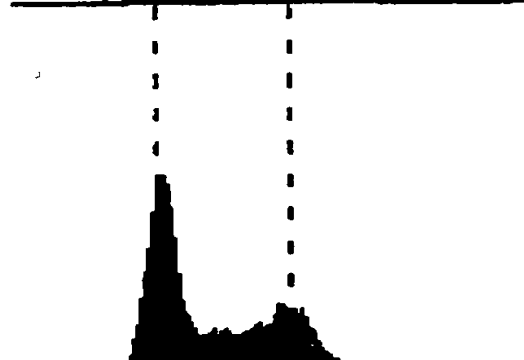
[Figure 13]



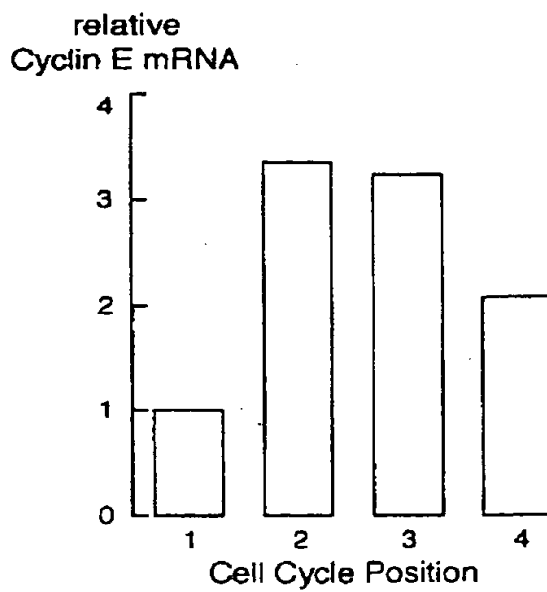
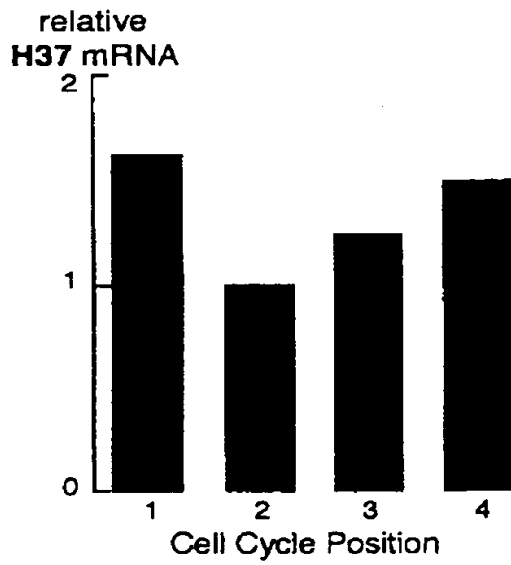
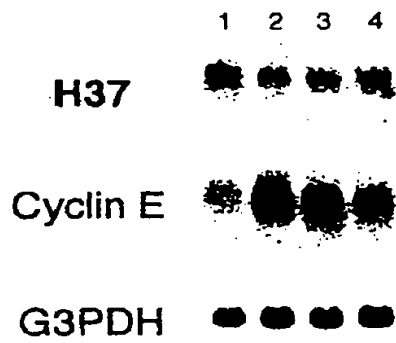
[Figure 14]



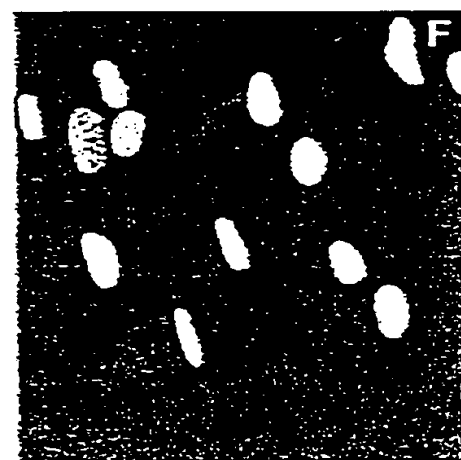
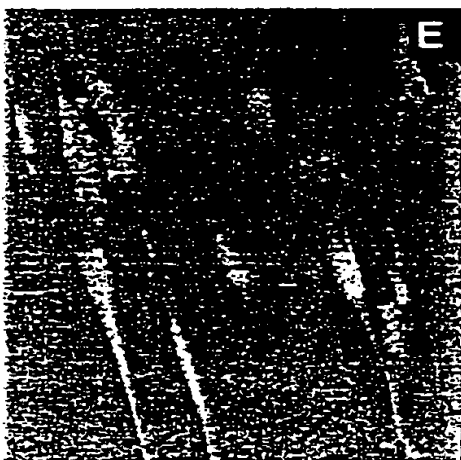
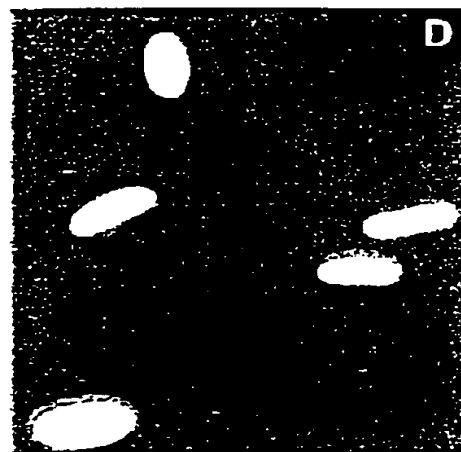
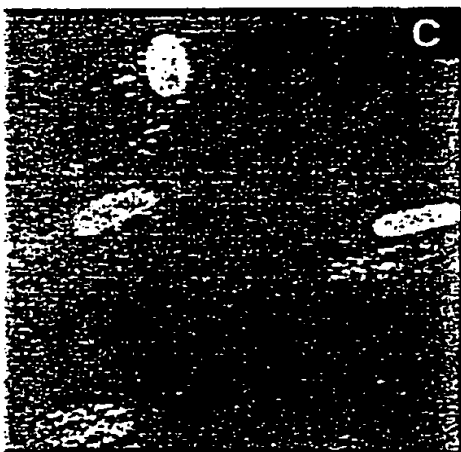
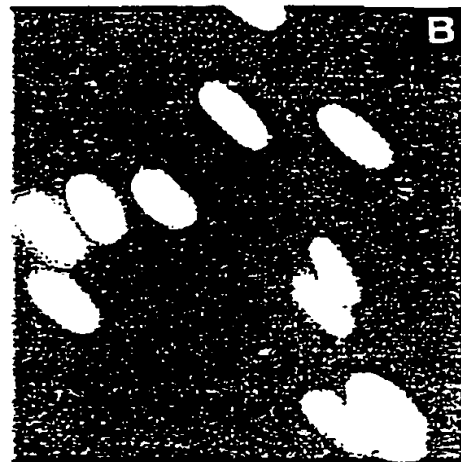
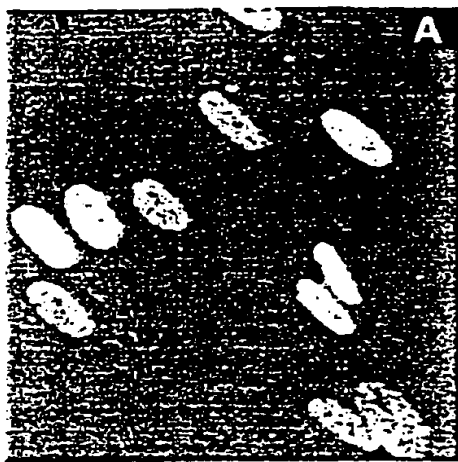
[Figure 15]

1 (G2/M)**2 (6h)****3 (12h)****4 (18h)**

[Figure 16]

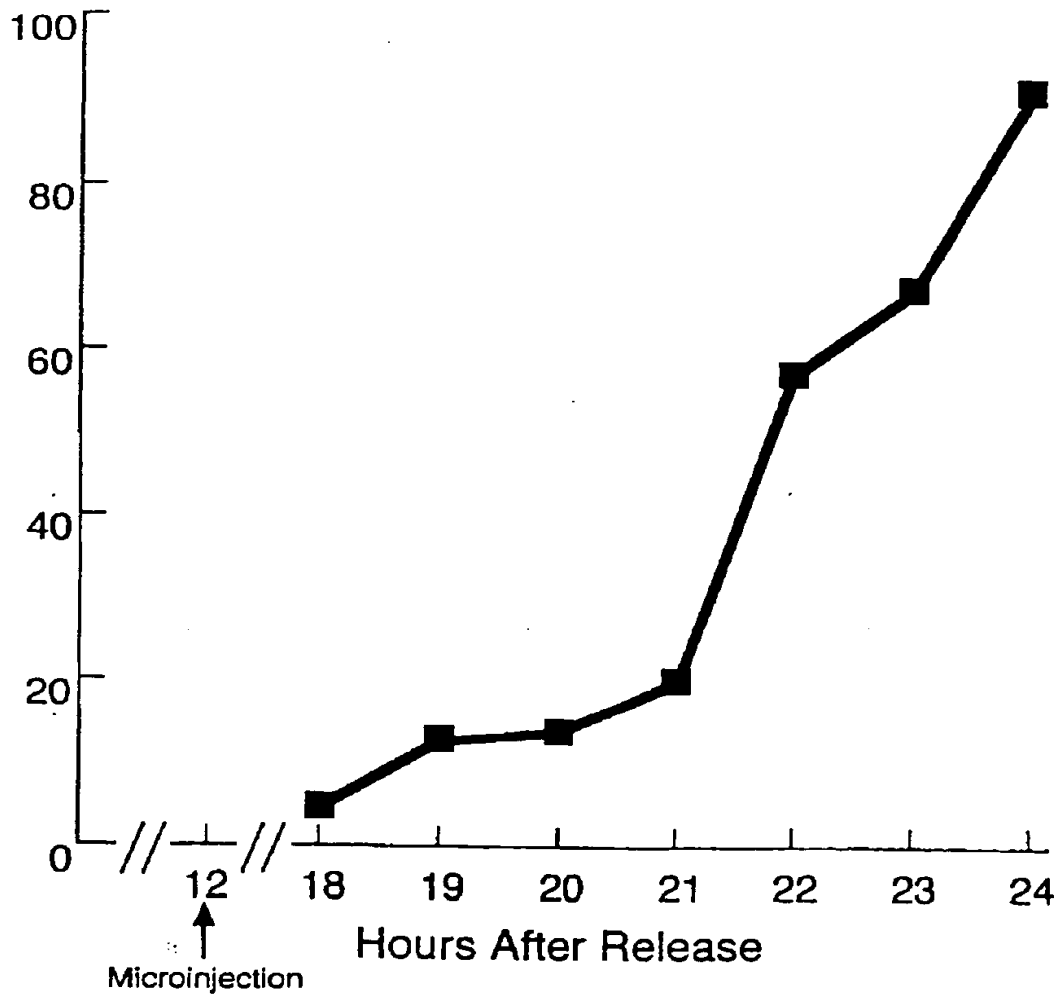


[Figure 17]

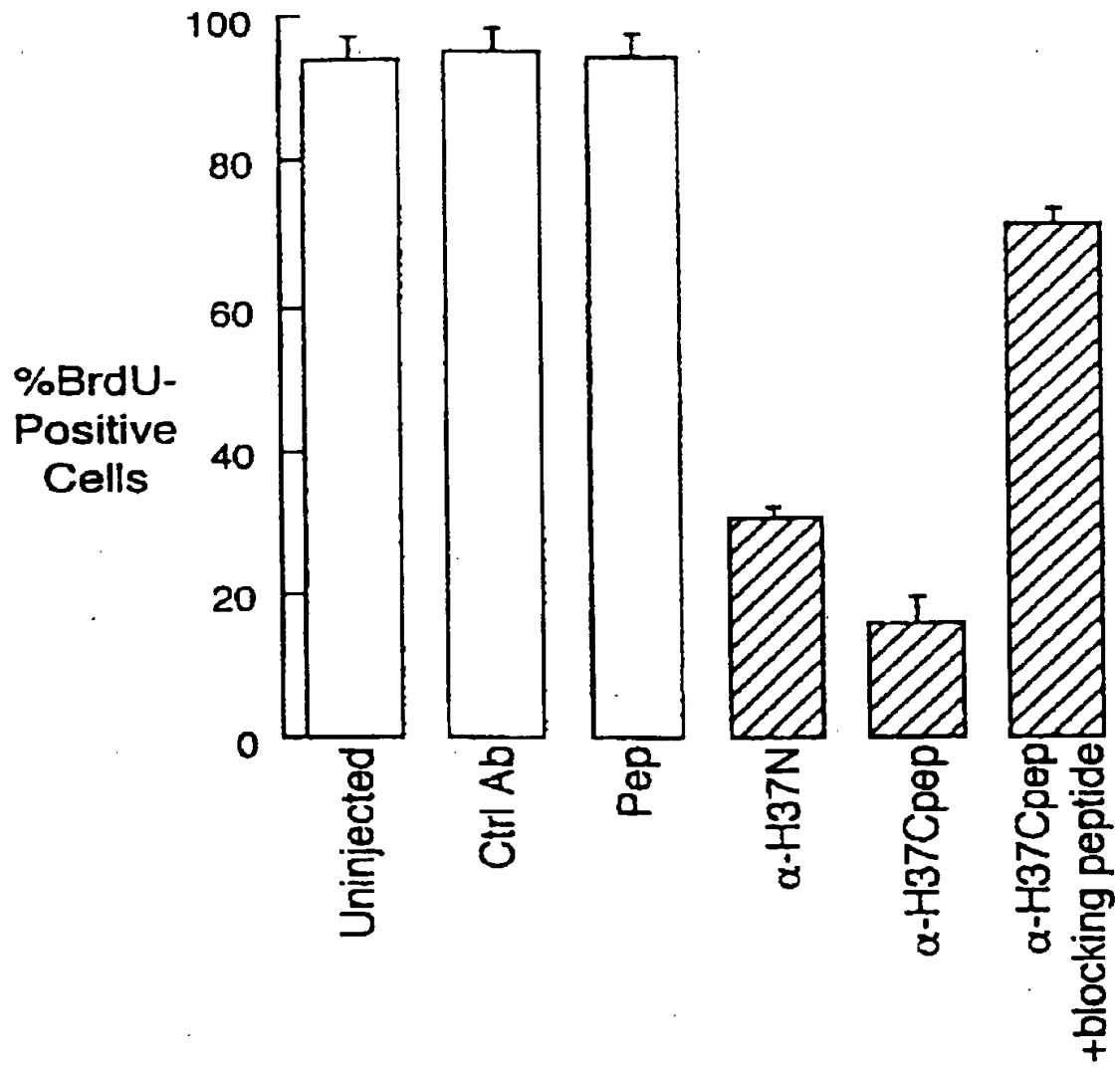


[Figure 18]

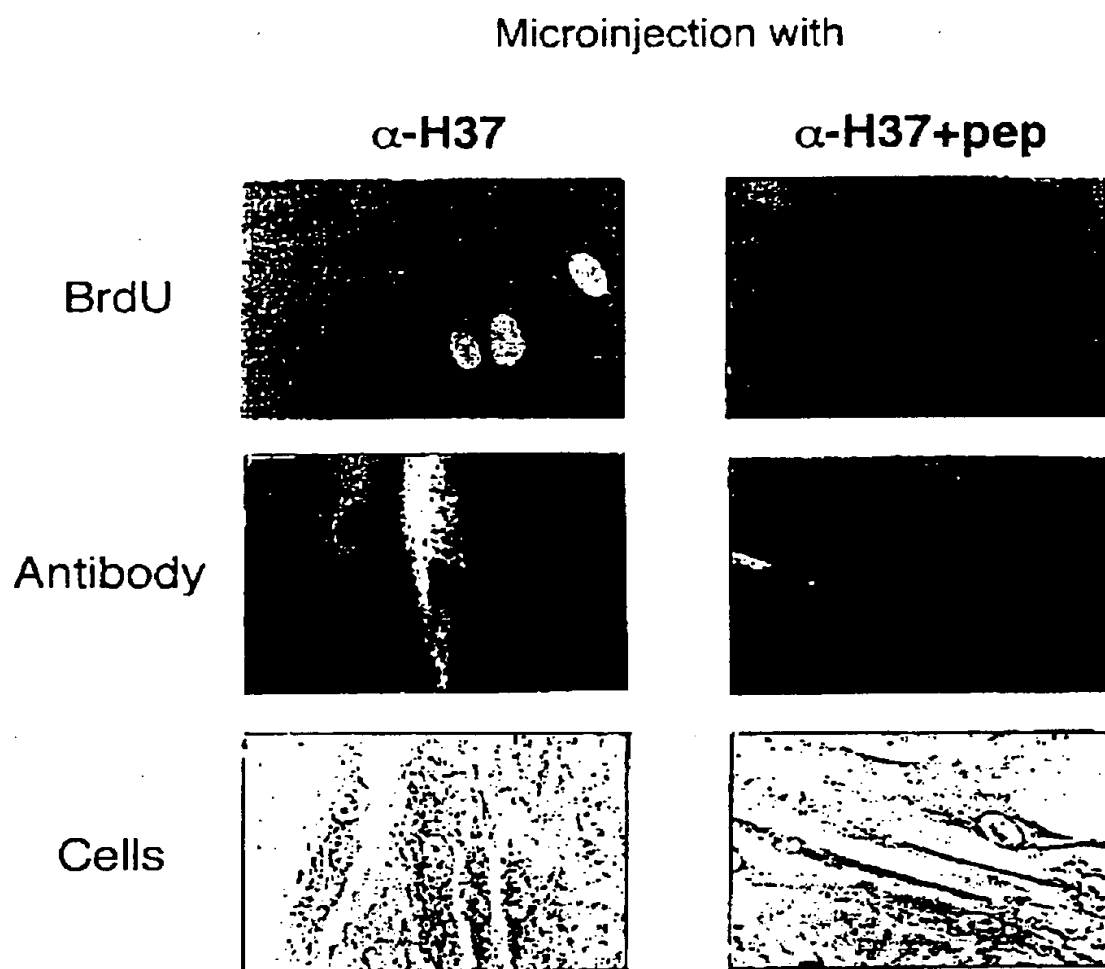
%BrdU-Positive
Cells



[Figure 19]



[Figure 20]



[Document Name] Abstract

[Abstract]

[Problems to Be Solved]

5 The present invention provides human H37 protein, a regulatory subunit for Cdc7 that regulate replication of human cells, human gene encoding this protein as well as cDNA thereof, an antibody against H37 protein, and a method for controlling the proliferation of human cells using those genetic engineering materials and antibody.

[Means to Solve the Problems]

10 This application provides human H37 protein having an amino acid sequence of SEQ ID NO: 1 or NO: 2; human gene encoding the protein; cDNA of the human gene which has a base sequence of SEQ ID NO: 3 or NO: 4; DNA fragment comprising a partial sequence of the cDNA; recombinant vector having the cDNA; antibody against human H37
15 protein; and a method for controlling the proliferation of cells by introducing the above DNA or antibody into the cells.

[Selected Drawing] None